tcatattttt	attgtacata	gacattgtgt	gtgtacttca	ttgtaggagc	aataaatcac	1500
ccattctgtt	taatgaagtc	atctgtgttt	ctgtagcact	ttctcaatgc	cttattatac	1560
ttccagtaat	tattgatgaa	cttcaacatt	atactacaat	aaatgtcttc	tcatttgctg	1620
tacagacagc	cagaaagtat	aagtaattag	aatttgttct	gcactctact	tttatcgtga	1680
aaaagtaatt	gaaagtaaat	aagatattta	ttcagaaaaag	ccactacctg	aatgtgagca	1740
tatctgaact	aataaaaata	gtatagttaa	tacagatata	tagcaagatg	tcctcaccat	1800
ctacatctgt	aataatatat	agtggtaatt	gactcccata	atggttatac	caaagccctg	1860
aaggattctt	aacttttact	ttcattttta	atgcaaaacc	aggttttcat	atattttgct	1920
tgcttaaagc	aagatatact	agctcttatt	tttcttacta	aattacatgc	atacatatat	1980
ttcaaatttt	attttagatt	atttgtgctc	aaccccatac	gttgtcatag	agtagtgctt	2040
ttcaactcgt	tttccaagga	tcactaattc	tgtagaatgt	taatagatgt	tatgtgaggg	2100
ggagaaagtc	attcagcttt	gtaaatgctg	aattaaatat	ttctttttga	acaaagctgg	2160
aggcatcatg	ctacccaact	tcaaactata	ctacagggct	acggtaacca	aaacatcatg	2220
gcactggtac	aaaaacagac	acatagacca	atggaacaga	atagagagcc	cagaaataaa	2280
accgcacacc	tacaactgtc	tgatttttaa	caaacctaac	aaaaacaaac	aataaggaaa	2340
ggattcccta	ttcaataaac	agtgctggga	taactggcta	gccatatgca	gaagattgaa	2400
actggacccc	ttccttatat	acaaaaatta	tctcaaggtg	aattaaagac	ttaaatgtaa	2460
aacccaaaac	tataaaaaacc	cacaaagaca	acctaggcaa	tacctttcag	gacataggca	2520
caggcaaaga	tttcatgaca	aagactccaa	aagcaattgc	aacagaagca	aaaatggaca	2580
aacggaagtt	aactaaacta	aagagctict	gcacagcaaa	agaaactatc	atgtttcttt	2640
tgagtaaaca	gcctacagaa	tgggagaaaa	tttttgcaaa	ctgtgcatcc	aacaaaggtc	2700
taatatggag	catctataag	aaacatacaa	atttacaagg			2740

<211> 2828

<212> DNA

<213≻ Homo sapiens

cttcatataa	gtgggatcac	gtaacattta	tccttttgtg	tctggcttat	ttcactttgc	60
ataatgitti	caaggttcat	ccatgttgca	gcatgcataa	aaacttcatt	ccttaaggtt	120
gtatattatt	ccattgtatg	catatgccac	attttgttta	tccattcatc	cattaatagg	180
catttgagct	tcttccacca	ccttttggct	attgtgaata	atgttgctat	ggactttaca	240
gtacaaatgt	ctgaccaagt	ctctgctttt	gattctttgg	ggtatataca	taaaagtgga	300
attgctggat	catctggtag	ttttacattt	cactttttt	agggactgcc	atattgtctt	360

```
ccacagtgac tgcaccattt tatactccct aaggcaatac gcaagggttc taatttctcc
                                                                     420
                                                                     480
atatteteae caacaettgt tttetgtttt gaggataata gecateetga gaggegtgat
                                                                     540
gtggtacctc attgaggttt caatttgcat ttccctaatg atttctgatg ttgagtgttt
                                                                     600
tttcatgtac ttcttggcca tttgtatatc tcccttgaag aaatatcagt atcaagtcat
ttgctcatgt ttaattgaac tgctttgttg tggtggtttt tactaagttt ttggagttct
                                                                     660
                                                                     720
ttaaatatcc tggatattaa tctcttacca gtatatgatt tacaaaagtt ttctcccatt
                                                                     780
ttgtaggttg cctttatctt ctagatagta tcctttgatg cacaaaaaat taattttgat
                                                                     840
gaaatccaat ttatctattt tttggcataa tatccaagaa atcattgcta aatccagtca
                                                                     900
ctaagetttt etegtgtgtt ttettetaag agttttatag ttllagetet tatgtttaag
tetteaatee attitgagti aactittgig eagggigtaa gaagtatite atgatticaa
                                                                     960
aattttgtgc gtggcataag gagtatttcc tgattttaaa gtatcaggaa atatttccaa
                                                                    1020
aagetgatac elecagetti attgitetti ticaagattg aettegetae teatggtett
                                                                    1080
tigiggitet atataaatii iggaatiitti titatiietg taaagaaagi geeatigaga
                                                                    1140
                                                                    1200
ttttgatagg gattgttttg aatgcataga tcactttggg tagtatgggc atcttaacaa
                                                                    1260
tattaagtet teeaattaat taacatggga tatettteea tttatttaca tettetttga
                                                                    1320
tttctttcag caatgttttg tagtttttag tgttcaaatc ttttaccctc ttggttaagt
                                                                    1380
gtattcctat tttattctct ttgatgttaa tgtaaacggt cctttcagat cgttcattgc
                                                                    1440
tagictataa aaatgcagct gittictigg tgiggigatt tigtaaccig caagittigca
                                                                    1500
cgattcattt attacttaac agtttacttg tgcagtcttt agaattttct acctgtaagt
                                                                    1560
ttacgicate tgtgaacaga gataatttta ettetteett tecaaittga atgeetttea
tttettttte ttgeetagtt gttetagett geetaattge tgtaggaett eeagteetgt
                                                                    1620
                                                                    1680
gttaaataga agtggtgaga gtgactgagt atgatgcatt ggattttgaa atagaaatag
acacttagta aaaaaaacaa aaactgggaa aattctaata aagcctatag tttagttaat
                                                                    1740
agtattatac cagigitcat cicilagitt tcataattat acigiagita titaagatti
                                                                    1800
                                                                    1860
taatgctaga ggaagcttgg taaagagtat atgggaaccc cgcactattt ttgtaactcc
tetgtaaate ataaatgate teaaaataag ttttttaata tetgaaggaa aataaaagea
                                                                    1920
                                                                    1980
tacagagtaa atacattggg agtcctgaaa tatctttcta ggcaaataag gcatactttc
                                                                    2040
actiticati agitaattac ccatagiaag tgiccaccac clcigccact ciggiatata
                                                                    2100
aagtoctaga gggcaactgg ccagacatca cattaaagaa gcactatggc actggactga
gicaaticig aaiticagci tiggggatti cicagggcaa agiigtitaa tatcicigaa
                                                                    2160
                                                                    2220
atatgaaaat atettteeag gattgttgtg aatattaaat aaggteatgg ttaaaaaage
actiaateca tggcaggtat ttaataaata teettietti teletigiti taaacatila
                                                                    2280
                                                                    2340
agactacaaa gggaaaacga gatatggtac ctttcacata agggccttac attctaaatt
gggacaaaat gtacacatat gttaaataat tatcaaataa tagctttaca tictattaca
                                                                    2400
                                                                    2460
agcaagtgct aagatccagt agcgagtggg acagtgtgca gcagagcaca gcacacaca
```

aagtagagaa	atgtgagctg	gcatcccgg	ggcagccttc	aaaggagtgg	ggctgtactg	2520
agcttggtgt	gcacagataa	gaaaatgttg	ggggtggtcg	ggtgcggtgg	ctcgcgcctg	2580
taateccaac	actttgggag	gccaaggtga	gtggatcacc	tgaggtcagg	cattcgagac	2640
cggcctggcc	aacatggtga	aaccccgtct	ctactaaaag	gtacagaaag	ttagccaggc	2700
ctggtggtgg	acacctgtag	tcccagctac	tcaggaggct	gaggcgggag	gatcgcttgg	2760
gcccagaagg	cagaggctac	agtgatccac	tgtactccag	cctgggcaac	agagcgagac	2820
cccatctc						2828

<211> 2735

<212> DNA

<213> Homo sapiens

attccacttc	cggtgtcttg	gtcttatcac	tgtaacttga	gctgagactt	tacttccatg	60
tctccccca	ccctcaggag	tcctcagggt	ttccactaaa	gcccctttc	attgagccta	120
caagccttgc	tgggaggcac	aggcagcgtc	caccccgatg	atgtcctccc	gccggcatcc	180
cgcagccttc	gccttcctgg	ttcacctgtc	gcttacctgt	ctcgctctct	cacctctgct	240
ggtcagcaga	tctgaatatc	tctttgtgag	gagtcagaga	aaggagatga	tgcaaagaga	300
aggaagcaaa	aacaaaacta	aaagaacttt	gaaggtatat	gagaagaaaa	agctgaaaag	360
aaaatgtctg	cctcactccc	tcaaaaattc	taccatagag	caaaggttga	cgtccagaaa	420
acctcaagtt	tatcagtccc	agatgctggc	agtagaagca	taaagctgag	gtgaattttc	480
cttactattt	tttccccaga	gtctagaaca	gtaactgcct	cccaatgtct	atcacaggga	540
ccagattttc	tccactacat	aggatgtgat	atgtaggcag	ttaaatggca	ttatagacac	600
aataagcaaa	attgaaacag	agacaccctg	ttggataaaa	tatttacatc	ggtgttctgt	660
ttccaaactg	tgggctgaga	cctataaatg	agtcatgaaa	tcaacctttg	tgggttgcaa	720
ccagcatttt	cacaagaagg	gattaagaaa	caacttcaga	atgtatcttg	tgtagtaaca	780
gtgactactg	ttttgcaaaa	tgtttgggtc	agttgtacac	atgcatattt	tacagggtga	840
tgtaaaaaaaa	aaatgtttt	tttgagacag	ggtctcactt	tgtctcccag	gctgaagtgc	900
agtggcaccg	tcatggctca	ctgcggcctc	aacctcctgg	gctcaagcaa	tcttcctgcc	960
ttagcctcct	gtatatctgg	gactacaggt	gtgcaccate	atgcccagct	atttatttca	1020
ttitttgta	gagccagggc	ctcactttgt	tacccagact	ggtctcgaac	tcctgggatc	1080
aagcaatcct	cccacatcag	cctcccaagg	tgctgggata	acaggcgtgg	gccaccgtgc	1140
ctggcctgaa	atgtcttttt	aactgtaggc	tggggtccgc	aacgtttgag	aagcactgaa	1200
atacagggca	atttattcac	tgtccaagaa	cattacttct	atgatgtctt	cagataaatt	1260 -

tgaaaatatt	tgtcttaccc	aaaattccag	ttacatatcc	attttcagga	tactctcttc	1320
tgagtgagat	aaactcaaca	ctggcaggta	tagcaagaaa	acaaaagact	ctgccaaata	1380
ttgctgcagg	gtgagaacca	ccagcacaat	gtggaaatcc	agtgtaggta	aagattcgcc	1440
ccaaatgtgt	gatcaggaaa	taaaacctgt	taggtgtcag	tgtcatgaag	ttcagtgtcc	1500
aaatgttgtc	tttcttctgg	taatggttac	ttttttttt	ttttcctgag	ataaggitta	1560
gcttttgtca	ctcaggctcg	agtgcaatgg	tgcgatcttg	gctcactgca	acctccacct	1620
cccgggttca	agcgattctc	ctgcctcagc	ctcccgagta	gttgggatta	caggcacccg	1680
ccaccacgtc	cagctaattt	ttgtatttt	agtagagatg	gggtttcact	atgttggcca	1740
ggctggtctg	gaactcctga	cctcaggtga	tccacttgcc	tcggccttcc	aaagtgctgg	1800
gattataggg	gtgagccacc	gcacctgggg	tggttactgt	ttaaaaaaaaa	agtcacctga	1860
ctcagaccca	ctgagcactt	ggagaacatc	tacaggtaga	aaacaaatgt	tacaaatatt	1920
taaatatcct	gatggttttg	gaagtcaagg	ggagagagat	cttacctctg	tctaccaccc	1980
aaccctctcc	accaaggtca	ctataaatac	aaagagcata	gctigggcia	caggaaagaa	2040
ggcattcata	tgtataaatg	tgtgtgtata	tatacactgt	tttttctttt	ttaagagact	2100
gggtctccct	ctgtcaccca	ggctggagca	cagtggtacg	atcatagete	actgcagcct	2160
ccaaaggtgc	ggctgaagtg	atcctcctac	ctcaacctcc	ctagtagctg	ggattaaaga	2220
tacacaccac	cacatggcta	agaaggcatt	ctagccctga	gtggtggtga	cacaggttat	2280
atacttgaag	tatacacaca	cacatatata	tgtatacacc	catatgtaaa	tataacacaa	2340
aaaactatgt	atttcactct	atgtatatat	atttaccatt	ttaaccgttt	ttaagtgtac	2400
aattcaatgg	cattaaatac	attgttgtgc	aaccaacatc	accatccatc	tccaggactt	2460
ttttggcatc	ccaaacggaa	actccatagc	cattaaacaa	caatttctca	tttccccttc	2520
cctccagcca	tgggcaacca	ccatgctact	ttctgactct	atgaatttca	ctactctggc	2580
tacctcatat	aagtggaact	gtactgtact	ggtcctttta	tgtctgactt	atttcactta	2640
gcataatgtt	tttgaaggtt	cgtctatgtt	gtagtatgtg	tcagaatttg	ctaccttttt	2700
taggctgaaa	aataaacgat	tcacaatata	ttttt			2735

<211> 1847

<212> DNA

<213> Homo sapiens

<400> 4642

gegtgegee tggggeea gttggggeg geegtgeta geggggga ageggggge 60 agtgeageea atgggaggeg gegetgeeta geggetggta ggeggtgeet gegegggtgt 120 taggttageg egaggegtga eetagttgae aggetetgag gtgetgetgt ggeggegtee 180

240 gcggggctga ggcgggtggg agccggagcc gagcgcgggc tgagggagga gggcggcgac 300 tggagagegg egageggega geagegeagg aegeagagee tettteaett ttteeetget 360 gagtgcccc tcccaccct cccactccac acacacctg tttgcccgtg agcctgggga 420 acttgeaget taaageeage caccecaeg geaacatgta ceccageaac aagaagaaaa aggtgtggag agaggagaaa gaacgtttat tgaagatgac cttagaagag agacgcaaag 480 540 aatacctaag agactatatt cccctgaaca gcattctatc atggaaggag gagatgaagg 600 gcaagggcca aaatgatgaa gaaaatattc aggaaacatc ccaggtgaag aaaagtttga 660 ctgaaaaagt ttctctctat agaggtgaca tcacattgct agaggtagat gctatagtca 720 atgccgcaaa tgccagtctt cttggaggag gaggtgtgga tggctgtatt catagagcag ccggcccctg tttgctagct gaatgtcgta acctgaatgg ctgtgatact ggacatgcaa 780 840 aaatcacatg tggctatgac cttcctgcaa aatatgtcat ccatactgta gggccaatag ccagggccat attaatggtt cccacaagga agaccttgca aattgctata aatcatctct 900 960 gaagetegtg aaagaaaata acateegate agttgeattt eeetgeatet caacaggeat 1020 ttatggettt eccaaegage etgetgeagt eattgeeete aaeaceatta aggaatgget tgccaagaat caccatgagg tggatcggat cattttctgt gtcttcttag aagttgactt 1080 caaaatctac aaaaagaaaa tgaatgagtt tttctccgta gacgataata atgaagaaga 1140 1200 agaggatgtt gaaatgaaag aagattcaga tgagaacggt ccagaggaga agcaaagtgt 1260 ggaagaaatg gaagagcaga gccaagatgc agatggtgtc aacactgtca ctgtgcccgg 1320 ccctgcttca gaagaggcag ttgaagactg taaagatgaa gattttgcaa aggatgaaaa 1380 tattacaaaa ggcggtgaag tgacagatca ttctgtgcgt gaccaagatc atcccgatgg 1440 acaagagaat gattcaacga agaatgaaat aaaaattgaa acagaatcgc agagctcata 1500 tatggaaaca gaagaacttt catcaaacca agaagatgcc gtgattgtgg agcaaccaga 1560 agtgaticca ttaacagagg accaagaaga aaaagaaggt gaaaaagctc caggcgagga 1620 cacacctagg atgcctggga aaagtgaagg clccagtgac ctagaaaata ctccaggtcc 1680 tgatgcaggg gcacaagatg aagcgaagga acaaagaaat ggaactaaat gacaatcctc 1740 agcategeaa ggeeteteet ggetetgggg gageteggga agatageage acaegetgtg 1800 gaggaggtg ggggtggggg gaaggcaagt cccatggaag gacggggaat cctttactct 1847 aatttctcca gctgcatttt gttccgttta tctgcagaaa aagaaag

<210> 4643

<211> 2987

<212> DNA

<213> Homo sapiens

gtttgtcacc	ttgtactcag	ccctgtgta	cagaacatgg	cacagaagcc	atttttgtgg	60
cagcggtggg	gattcggccg	tctcaccatg	ctgctgcaca	gtcttagagt	ccttgctcag	120
agccgggctg	tcgtttattt	aacatccaac	ggtcggtgtg	tgtgtgtgtg	tgttgtgggg	180
gagtgggggg	gtgttctctt	cctacaatgc	tgcctcccag	gttcctgcag	cagccacaca	240
tatctctggg	tggctgggga	ggaggaccag	gtcagaatgc	agtaggacaa	ggttcttttc	300
tccaggaggc	ccagagagga	tgatgggtca	ggtaaaaggc	cctgcactta	actctctgcc	360
cctgggagac	ttatttctcc	tctggggact	tgtttctgca	tccacaggtg	aaagtggttt	420
aaatgagaag	ctcttaaact	ttcttggttt	ggcagcagag	ttatttttct	agacaaagta	480
taatgtgaaa	gtttaaatgg	atgaaaatga	actgaggagg	ccccttggt	gggagccact	540
ggggtgtgtg	tggcagttgg	gggatcctgc	atggtctccc	ttcattcccc	taatgcctct	600
tgaggcccct	cagtgtgggc	cattaaaagg	cccctagacc	agcagatccc	ttctacagat	660
ctgcttattg	aaaaacctgc	tctgtgccag	gcagtgcctg	gaattctaaa	cctgaagctc	720
ccaaatgctg	ctctccgggg	cctggctcac	attatcaagt	cacttgggaa	gcttaacaaa	780
taacgtgggt	tigiggtage	tectigacag	gtgtgcttat	ccgtgtcgcc	tgtcttggct	840
tccgatgcaa	agccagcttg	ggaacccctg	gcctcatctc	cttcagagtc	ttcgtttgac	900
cttggtgggg	catgaggacc	tgctggccct	tgttaaagcc	agtgaaggtg	gtgaccctcc	960
cgggccatct	gttaagttgc	cctgttgatg	gggttgggag	gagttagtag	agaagcaggg	1020
cttcttttgc	ttagcagccc	ctgcccagac	ccatctgtgt	gtctctcaga	caagcctgtg	1080
ccggaggagt	cggagggccc	tggcagccct	ccccctaca	agatgatcca	gaccattggg	1140
ttgtcggtgg	gtgccgctgt	ggcctacatc	attgccgtgc	tgggcctcat	gttctactgc	1200
aagaagcgct	gcaaagccaa	gcggctgcag	aagcagcccg	agggcgagga	gccagagatg	1260
gaatgcctca	acggtgggcc	tttgcagaac	gggcagccct	cagcagagat	ccaagaagaa	1320
gtggccttga	ccagcttggg	ctccggcccc	gcggccacca	acaaacgcca	cagcacaagt	1380
gataagatgc	acttcccacg	gtctagcctg	cageceatea	ccacgctggg	tatgttgcct	1440
tgactacagc	tgcccctgcc	taactataag	tcacccgctg	tgtcttgagg	ctgggtacgg	1500
gctagtgttc	tgaagtccct	gaaacctaga	atgttagagg	tggagggaag	cataggtaga	1560
gatctagccc	agtgtggccc	cagagectgt	tattcactgc	ccttcgtgca	gattttgtca	1620
ttcctgtgta	ctacctgtag	tattatttac	ttgatgtttt	atttaaattg	actctttaaa	1680
aaaaaaaact	aagttaccat	tgttctagga	aaaaatatga	gaaatcatgg	gtttgtcata	1740
gttacgtttt	ttcctagcat	acgttaagac	atgtttacca	tgagccacca	aaaattctca	1800
tgcgagtcac	cctttggaaa	atgctgaata	ttagtcccac	ccttttattt	tgttgctgca	1860
gacactgaag	cctaaggagg	gagggggtca	cagcagaacc	gggtctcgtg	cccagcccag	1920
gtgggtgggt	ccccactgtg	ggagaggcta	ggcccctccc	ccaggtcagg	agcigtcicg	1980
gcctgggtga	aggtggctgg	ctgactcaga	ctgtacccac	agggaagagt	gagtttgggg	2040
aggtgttcct	ggcaaaggct	cagggcttgg	aggagggagt	ggcagagacc	ctggtacttg	2100
tgaagagcct	gcagagcaag	gatgagcagc	agcagctgga	cttccggagg	gagttggaga	2160

tgtttgggaa	gctgaaccac	gccaacgtgg	tgcggctcct	ggggctgtgc	cgggaggctg	2220
agccccacta	catggtgctg	gaatatgtgg	atctggtatg	ctgttggcag	gggacgtggg	2280
ggtctcgggt	agggagggca	gtgtcctaca	aaggtgggag	tcagtggttg	gagcaccgtg	2340
aaaaccttgt	ctcgtgcagt	ctcagccgag	acctcacctg	cctgctgtta	cactttgccc	2400
accttatgat	gctcagcttc	tgccttccct	gctcttggat	gcctgctttc	cctttatcag	2460
cacccagtct	ttccatctgt	atctcagggg	cgttggtcct	aagtttttct	gggacttctc	2520
actgtcacac	tgcagggatg	ccttccaggt	catcttctct	gactcttgtt	ttctttggtt	2580
tgttatttt	tectecceaa	aagcttgtcc	ccacttgaga	ataaatatat	cagaacccac	2640
tttgcctccc	tccttcctag	caccacttgt	gacatgagtg	ttaccccagc	caccagcctt	2700
ctcctctcca	ctcagggtgt	cattcccaga	gctgcgcctt	cctcttcatt	ccccagctgg	2760
tccttgccct	caccccatgg	tcctcaaact	gctgcttctc	aaatccctag	ttctggggca	2820
agagcgggca	ggagtcctat	cagtctttcc	attaacaagt	acttacgcgg	ctgggcacag	2880
tggctcacac	ttgtaatccc	agcactttgg	gagactgagg	caggaggatc	gcctgaggcc	2940
aggagtccga	gaccagcctg	ggcaacatag	caagactctg	tctctac		2987

<211> 2253

<212> DNA

<213> Homo sapiens

aatttigcaa	cgacgggcgc	ggaggaggag	gttcccggaa	gccacgcgca	gctggagcag	60
cggcgaccgc	agctggaggc	ccggagcgcc	tgcggggctg	gcagaggcga	gggaggttgc	120
gggtagggag	ggcggactgc	gcgcgccccc	tgcgtcccgc	gcacctcggg	gccggtccat	180
gctcccgacg	gctgcgggct	tcagcatctg	gggccaggtt	ggggcagcaa	gggaggcgcc	240
gcgatgccag	acgaaaatat	cttcctgttc	gtgcccaacc	tcatcggtga	gtgctgccca	300
cggccccggg	ccgaacgaga	gggcgagggg	ccttggcggg	ctccctgatc	ctgccctgtc	360
tccccgcgtc	ccgttccaag	gttatgcccg	gattgtcttc	gccatcattt	ctttctactt	420
catgccctgc	tgcccctca	cggcctcctc	cttctacctg	ctcagcggcc	tgctggacgc	480
tttcgatgga	cacgetgete	gcgctcttaa	tcaaggaacc	cggtttgggg	ccatgctgga	540
catgetgaeg	gacegetget	ccaccatgtg	cctgttggtc	aacctggccc	tgctgtaccc	600
tggagccacg	cigitatic	aaatcagcat	gagtttggat	gtggccagtc	actggctgca	660
cctccacagg	tctgctgcga	ttctgggggc	ctgggccact	tggagacatt	acagtggggt	720
gggctgagcc	tagatggaag	ggtgtctggg	atggcactag	gcatcctgga	agggctcctt	780
tttagctctg	ctttgtagaa	agaccccagg	caaacctgcc	tctctttaaa	gcttcagtca	840

tgtttacagt	tgcattaagc	aattcttagg	actcttctgg	ctgggtattc	taggattcca	900
tataggtgac	agcttatagg	aaggggcatg	ggagctttca	aatgtggcat	ttgaatcctg	960
actctgcccc	caaataggta	actgtggctt	tcggcaggat	actggtcctt	agccgtgccc	1020
agttttccat	ctgtacaata	ggtgttaaca	tgaggattgg	atgcatggta	gtggtcatga	1080
aaccttgtgg	gttcccctgt	tttctcccca	agttctgtgg	tccgaggcag	tgagagtcac	1140
aagatgatcg	acttgtccgg	gaatccggtg	cttcggatct	actacacctc	gaggcctgct	1200
ctgttcacct	tgtgtgctgg	gaatgagctc	ttctactacc	tcctctacct	gttccatttc	1260
tctgagggac	ctttagttgg	ctctgtggga	ctgttccgga	tgggcctctg	ggtcactgcc	1320
cccatcgcct	tgctgaagtc	gctcatcagc	gtcatccacc	tgatcacggc	cgcccgcaac	1380
atggctgccc	tggacgcagc	agaccgcgcc	aagaagaagt	gacgctggag	ccccgggtcc	1440
tggctgccca	cctgccctgg	gagtcttgct	gtgccacaca	gctccccacc	ccctgctagg	1500
aggtcccagt	ctcacgcctt	cctcatgtgt	tgttctacct	gctgggatgg	gggtcagcct	1560
ctctttggtg	acgtcacgtt	ctctgggatc	ctgaggaccc	gggcctcaaa	tcagggagga	1620
tacgcgggag	gececeteca	tecaggeggt	gctcctgggg	tgccgggacc	gggcagtgtc	1680
acaccctgcc	tgctcagtcc	tggggtccga	gatgctaggg	acgcttgagt	gagggaggtg	1740
gtgtgagggc	caggtttcct	gaaaggcggg	agtcagacct	ccgccccag	ccagagcaag	1800
cttggggcac	catgcccagg	agggaagaag	ccatccacag	ccttccctgt	caccggctcc	1860
tctgtcctgc	ctaccctggt	cctggcggga	cttcactatt	tgacttggtt	tcctttcaga	1920
tattcttggc	tcagggcctg	ggttgaggga	gcttagggaa	ggacgtccgt	ctgggtgctt	1980
ttcctccagt	ttgctggctg	gcttctccgt	ctacccacag	tgacctcaca	gagaggccct	2040
cctgccaccc	atgctcatgt	ggtgtcccca	ccgcccactt	gtttgatgtc	actgactgtc	2100
tacatglatt	tatattettg	atattttcta	ccctcactag	aatgtaaact	ccatgaaggc	2160
acagactttt	cttgttctct	tetgtatece	tagagtaaga	ccaacttgaa	cctggcatat	2220
agtagctgct	taataaatac	tegtetgtea	atg			2253

<211> 3001

<212> DNA

<213≻ Homo sapiens

aaaaaaaaaa	aaaaaaaaaa	aaaacagaga	agtcaaatgg	tttttggaa	tatggtggcc	60
ctccgtaccc	atcggttcca	catgtgtggt	ttcagccaac	tatgtattga	aaataaaatt	120
gcatccttac	aaatatgcag	acttttttc	cttgtcattg	ttcccttaac	aatacagtgt	180
aacagctatt	tacgtagcat	ttacattgta	ttaggtacta	tgagtcatcc	tggagttgct	240

```
300
gtaaaactta aacgtaaaac ttgaaatgag gatgatttaa agtatagagg aggatgtgca
                                                                     360
taggttatat gcaaatactc tcccatttta tattagggac ttgagcatcc acggattttg
                                                                     420
gtatccgtgg gggtcctgga cccaacctgc cacggatacg cagggacgac tgtatttggc
                                                                     480
atagaggeet tietaatgea ettaecaagg acaeetgeag eaggetgiga teeetgagae
                                                                     540
gagcgtgagt tlagtgaggg cagagttttt gttctggttc tcagctgtgt cccagcacct
                                                                     600
gggattgtcc ctggcacaca gtagctgctt agaaaagatt tgatgagagg gtgaccatac
                                                                     660
atgaggggaa attitectea gtateaaaac atgitgaaac teacacacti eeaagatgae
                                                                     720
gtattctcag gtctcctgcc gttgccgttc tctgccgtat gtgatggtgt tctgtcttac
cagaggetgt ggccctgaga aagatcacgt ctacctgcag ctgcaccacc tacctccaga
                                                                     780
                                                                     840
gcagctggcc acgcgcctgc ctggcatttc agagacagcc atgatcttcg ctggcgtgga
                                                                     900
cgtcacgaag gagccgatcc ctgtcctccc caccgtgcat tataacatgg gcggcattcc
                                                                     960
caccaactac aaggggcagg tgatggtgct ggctcctccc ccacagctgg aaagaaggct
                                                                    1020
gggacgacgg ggcccacctc gcagttgtct ctttagatct tacaggaaaa gatagatgtt
                                                                    1080
teetteaaga aagtaetgta itgiitteta gattgeacti taaattieta tiaceggagg
                                                                    1140
atggagggg citaalaati taliccicci tagtaaatig tcagagatac alcattigca
gettitteea tittgiaati aetiteetat atgatettgi gitattieta atgatettae
                                                                    1200
                                                                    1260
acatcaaggg atctttataa ttcattcctt tgagtggttt gtggttcaca cagagcttgt
                                                                    1320
cagtcactta ggctccttgt tgggcgaggt gggtggaagc tgttgctctc cctgcgtaga
                                                                    1380
cgaagaggtg aacggggtag aacagtctgg aacatcagtc tecectgctg atgtteetee
acctgccgtg ctcctgggtc tgagccggag cacaggtggt gagggccccg ggaacatggg
                                                                    1440
acacgggga cagtcgcaga tgctgacatt ggaggtcctc tgacctgctt gtaacagcag
                                                                    1500
gtgclcaggg gcagagggga aaclggggga taccllcgga agcttccctc tgaagaagag
                                                                    1560
                                                                    1620
tagetatggt cettactice etettagata eggtetitae ticceletet tittittiet
                                                                    1680
tggagatgea gicteaciet gilgetiegg eiggagagea giggigegat eicageteac
                                                                    1740
tgcaacetet geeteecagg ticaagtgal titteegeet eageeteeet agtagetggg
                                                                    1800
attacaggca cccgccatta tgccccgcta attittgtat tittagtaga gatggggttt
                                                                    1860
caccgtgtta gccagacagg tcttgaaccc ctgacctcag gtgatcaccc acctcagcct
                                                                    1920
cccaaagtgc tgagattaca ggcgtgagcc accaegcctg gcctacttcc ctetetetga
                                                                    1980
cetgeageae agacaecetg tiggggaagg tgggetggtg gaggeatggg cacettgaea
                                                                    2040
tttcacciga aaicticcii iccacaggic cigaggcacg igaalggcca ggalcagati
                                                                    2100
gtgcccggcc tgtacgcctg tggggaggcc gcctgtgcct cggtacatgg tgccaaccgc
                                                                    2160
cteggggeaa actegetett ggacetggtt geetttggte gggeatgtge eeegageate
                                                                    2220
gaagagteat geaggeetgg agataaagte eetceaatta aaccaaaege tggggaagaa
                                                                    2280
teigicaiga aiciigacaa aiigagalli geigaiggaa gealaagaac aleggaacig
                                                                    2340
egacteagea tgeagaagte aatgeaaaat catgetgeeg tgtteegtgt gggaagegtg
                                                                    2400
tigcaagaag giigigggaa aaicagcaag cictaiggag acctaaagca ccigaagacg
```

ttcgaccggg	gaatggtctg	gaacacggac	ctggtggaga	ccctggagct	gcagaacctg	2460
atgctgtgtg	cgctgcagac	catctacgga	gcagaggcac	ggaaggagtc	acggggcgcg	2520
catgccaggg	aagactacaa	ggtgcggatt	gatgagtacg	attactccaa	gcccatccag	2580
gggcaacaga	agaagccctt	tgaggagcac	tggaggaagc	acaccctgtc	ctatgtggac	2640
gttggcactg	ggaaggtcac	tctggaatat	agacccgtga	tcgacaaaac	tttgaacgag	2700
gctgactgtg	ccaccgtccc	gccagccatt	cgctcctact	gatgagacaa	gatgtggtga	2760
tgacagaatc	agcttttgta	attatgtata	atageteatg	catgtgtcca	tgtcataact	2820
gtcttcatac	gcttctgcac	tctggggaag	aaggagtaca	ttgaagggag	attggcacct	2880
agtggctggg	agcttgccag	gaacccagtg	gccagggagc	gtggcactta	cctttgtccc	2940
ttgcttcatt	cttgtgagat	gataaaactg	ggcacagctc	ttaaataaaa	tataaatgaa	3000
С						3001

〈211〉 3398

<212> DNA

<213> Homo sapiens

60	atttggacag	tggcccgtta	cagaatgaaa	ccaagaggta	tggcggctat	aaaatcttag
120	cctgctttag	aactgaagcc	aaaaggcatt	tgaaacctga	acaggctttt	acacacaaga
180	ggagttgcta	tgaaagccag	tgttigtcca	ccgtttcacc	tatctcaaag	ccctccctaa
240	ttatctaaga	agtagectat	ggagacgccc	ttaggaccct	tactcagacc	aaggggtgct
300	tagccgctac	agccaagcag	gttgtctgcg	ggatggccaa	tgagacctct	aactggatcc
360	cccttaaggc	caaaatttaa	aactctgggc	ctgataagtt	gtccaagagg	agcaagccta
420	aaatgctcgc	aatggatgtc	gcttctggcg	tacttcaagi	atagagactt	tecteatgee
480	ccacacggtg	actttctctt	cagtaaactg	actgttagat	atcagagttt	atcttgcaat
540	aaactggccc	gaaactaccg	ggaactgttg	atgactgica	gctactatcc	tttaaatcca
600	cagacggtag	accgtgttca	ggcggacgcc	ccctagagaa	caagatgtgc	atccgatett
660	agacagatgt	gttaccacgg	cggtgcagct	tacgaaaggc	gagcagggag	cagctttctc
720	tgategeect	aaagctgaat	ctcagegeaa	cagcaaacac	caggetttac	gttgtgggct
780	ggtacgcctt	actgacagca	taacgttaac	gtaaggatat	ctccgatggg	cactcaggct
840	cctcagcaga	aggetactea	ccaggagcgc	gagecatetg	catgtacatg	tgctacggtg
900	gcagctctct	aacagatcaa	ttcaaagcct	atccccgtc	aaaaacaaga	aaaggctatc
960	ccaggccacc	taaacccagc	aacaaggtée	gtaaatgcca	ctgcgaccag	ggggaacaac

gtctccgaag	aaacttgcca	ggagagaagt	gggaaattga	ctttacaaaa	gtaaaaccac	1020
accaggctgg	gtacaaatac	cttctagtac	tagtagacac	cttctctgga	tggactgagg	1080
catttgctac	caaaaacgaa	accgccaact	tggtagttaa	gtttttactc	aatgaaatca	1140
tecetegata	tgggctgcct	gctgccatag	ggtctgataa	tggaccggcc	ttcacctcgt	1200
ctatagttct	gtcggtcagt	aaggcgttaa	acattcaata	gaagctccat	tgtgcctatc	1260
gagcccagag	ctctgggcag	gtagaacgca	tgaaccacac	cctaaaaaaac	actcttacaa	1320
aattaatctt	agaaacttgt	gtaaattggg	taagtctcct	tcctttagcc	ctacttagag	1380
taaggtgcac	cccttactag	gctaggttct	caccttttga	aatcatgtat	gggagggtgc	1440
cgcctatctt	gcctaagcta	agagatgccc	aattggcaga	aatatcacaa	aagaatttat	1500
tacagtacct	acagtctccc	caacaggtac	aagatatcat	cctgccactt	gttcaaggaa	1560
cccatccgaa	tccaattcct	gaccagacag	ggtcctgcca	ttcgttccag	ccaggagacc	1620
tagtgtttgt	taaaaagttc	cagaaagaag	cactcactcc	tgcttagaaa	agacctcaca	1680
ccatcatcct	cacaacgcca	atggctctga	aggtggacgg	cattcctgct	tagattcatc	1740
actcccgcat	caaaaaggcc	aacagagccc	aactagaaac	atgggtcccc	aggcctgggt	1800
caggcccctt	taaaactgca	cctaagtcag	gtgaagccat	tagattaatt	ctttttatct	1860
acctcacttg	tttgtttttg	cctgttatgt	cctttgcgcc	ttcctactcc	tttctcctca	1920
cctctttcac	aacaggatgt	gtatttgcaa	acaccacttg	gagggccagt	acctccaagg	1980
aagtctcctt	tgcagttgat	ttatgtgtac	tgttcccaga	gccageccgt	accegegaag	2040
agcaacacaa	tctgccggtc	ccaaaagcag	gaagtgtcaa	ccttgcagca	ggatttggac	2100
actccaggag	ccaaactgga	tgtggaagct	ccaaagttgc	agaaaaagga	ctccaaaatg	2160
ttgactttta	cctctgtcct	gggaatcacc	ctgacactag	ctgtcgagat	acttaccagt	2220
ttttctgccc	tgattggaca	tgtgtaactt	tagccaccta	ctctggggga	tcaaccagat	2280
cttcattcca	taactcgtgc	ttctcgtcct	aaattatgta	ctagaaaaaa	tigiaateet	2340
cttactataa	ctgtccatga	ccctaattca	actcagtagt	attatggcat	gtcatgggaa	2400
ttaagatttt	atateceagg	atttgatgtt	gggactatgt	tcaccatcca	aaaaateetg	2460
gtctcatgga	gcccacccaa	gccaatcggg	cctttaactg	atctaggtga	ccctatgttc	2520
cagaaacccc	ctaacaaagt	tgatttaact	gttcctccac	cattcttagt	cataaaagat	2580
acactccaaa	agttcgagaa	aatctagata	ggcgccaaca	ggaacgagaa	aacaacatcc	2640
cctggtatca	aagcatgttc	aactggaacc	cctggctaac	tactttaatc	actgggttag	2700
ctggacctct	cctcatcata	ctattaagtt	taatttttgg	gccttgtata	tgaaattggt	2760
ttcttaattt	tgtaaaagaa	tgcatagctt	ctgtcaaact	tatgtatett	aatactcaat	2820
ataaccccct	tgttataatt	gaggaatcaa	cgatttgatt	ссссаааааас	acaagtgggg	2880
aatgtaatgc	ccaaccttgt	ttttactaac	cctgttttta	gactetecet	tttcctttaa	2940
ttacccagcc	ttgtttccac	ctgaattgac	tctcccttag	ctaagagagc	cagacagact	3000
ccatcttggc	tctttcactg	gcagccgctt	cctcaaggac	ttaacttgtg	caagetgact	3060
cccagcacat	ccaagaatgc	aattaactga	taagatactg	tggcaagcta	tatctgcaat	3120

tcccaggaat	tcatctgatt	gataacgccc	aaagcccccg	${\tt gtctatcacc}$	ttgtaatagt	3180
cttaaagccc	ctgcacctgg	aactgtttaa	tttcctgtaa	ccatttatcc	ttttaacttt	3240
ttgcctactt	tatttctgta	aaattgtttt	aactagaccc	tccctcccc	tttctaaacc	3300
aaagtataaa	agaaaatcta	gcccttctt	cgggccgaga	gaattttgag	cgttagccgt	3360
ctctcggtcg	ccagctaata	aatggactct	taattcgt			3398

<210> 4647 <211> 3818 <212> DNA

<213> Homo sapiens

<400> 4647

60 aagtgttage tegaagaaaa tatgeettge tacctagtte tagtagttee agtgagaatg acctcagcag tgaatcctct tctagctcat caactgaagg agaagaagat ttgtttgttt 120 180 ctgccagtga aaaccaccaa aacaatccag ctgttccctc aggaagtatt gatgaagatg 240 ttgtggtgat agaagcticc tccacicccc aggitacigc caaigaagaa aitaaigita 300 cctcaactga cagtgaagtg gagattgtaa cagttggaga aagctatcgg tctcgttcaa 360 cccttggaca ctccagatct cattggagcc agggttcaag ttctcatgca agtcggccac 420 aggagccacg gaaccgcagt aggatttcta ctgttataca gcccttgagg cagaatgcag 480 cagaagttgt ggaccttacc gttgatgaag atgaacctac tgtagtacca accacttctg 540 caagaatgga atcacaagct actagcgctt ccattaacaa ttcaaatcca tctaccictg 600 agcaggeete tgatactget teagetgtea ecagtageea acettecaca gtgteagaga 660 cttcagctac tettacaage aatagtacea etggeaette tataggagat gactcaagga 720 gaactacate tagtgetgta aeggaaactg geeeteetge aatgeeaagg ttacetteet 780 getgteecca geacteacea tgtggagggt egteacagaa ceaceatgea ttaggacate 840 ctcatacaag ttgctttcag cagcatggtc accattttca acatcatcac caccaccacc 900 atacteccea eccageigte ecaglitete effectitag tgatectget igecetgigg 960 aaagacctee acaagtacaa geacettgtg gagcaaatag tagttetggt accagetate 1020 atgaacagca ggcattgcca gtagacctga graaragtgg tatcagaagt catggaagtg 1080 geagittica tggageatet geattigace eetgetgeee tgittettee teeegageig 1140 caatettigg ceateaggee geigeigeig eeccaagica accittatea teaatagaig 1200 getatggate aageatggtt gegeageeee ageeceagee eeeteeaeag eeetetetet 1260 catcatgicg acattacatg ccacccctt atgcctcttt gacaaggcca citcatcate aagettetge etgeeegeat teteatggaa acceeetee teagacteag eeteegeete 1320 1380 aagtggatta tgttatteet cateetgtae atgettteea tteleaaata tetteleatg

caacatctca	tcctgtggca	ccccaccac	caactcactt	agccagtaca	gctgcaccaa	1440
tccctcagca	tcttcctcct	acacaccagc	caatttcgca	ccatattcca	gccacagcac	1500
ctccagcaca	gagactgcat	cctcatgaag	tgatgcagag	gatggaagtt	caaaggagga	1560
ggatgatgca	gcatccaact	ggtctttttg	tgttctgtgt	ttccaggcgg	gcacatgaac	1620
gcccccacc	ccatccacat	aggatgcacc	caaactatgg	tcatgggcat	catattcatg	1680
tgcctcagac	tatgtcctca	catcctcgac	aggctccaga	gaggtctgcc	tgggaactgg	1740
gaattgaagc	tggagtgact	gcagctactt	atacacctgg	tgcattgcat	cctcacttgg	1800
cccattatca	cgcacctcct	cgacttcatc	acttacaatt	aggagctctt	cctttaatgg	1860
ttcctgatat	ggcaggctat	cctcacatcc	gttacatttc	atcaggattg	gatggaacat	1920
cattcagagg	tcctttcagg	ggcaattttg	aggaactgat	tcatttggaa	gaaagattag	1980
gcaatgtcaa	tcgtggagca	tcccagggga	caattgaaag	atgtacatat	ccacataaat	2040
acaaaaaggt	aacaactgat	tggttctcac	agaggaaact	gcactgcaaa	caagatgggg	2100
aagaagggac	tgaggaagac	acagaggaaa	aatgtactat	ctgtttgtct	attttagagg	2160
aaggtgaaga	tgtgagacgt	cttccatgta	tgcacctttt	ccaccaagtg	tgtgttgacc	2220
aatggttgat	taccaataag	aagtgcccca	tatgcagagt	ggacattgag	gcccagctgc	2280
caagtgaaag	ttgacaccat	gtttcagaac	tcttgccctc	cctctcattc	ccatccttcc	2340
tggtactgca	gtcaaccaaa	gatggcatga	cttacctgcg	cagatttgga	agcattgaac	2400
ttagagtgct	ggctctgcta	tatggtacaa	ctaatgctag	acctacagtt	tatgtataca	2460
gttgattttg	atgtatttat	aaaagctttt	ttttctagat	ttgacatttt	tctgtatcat	2520
tttactgtat	ttttgcatgg	ttccttgtat	tgcatttctt	tgcacatatt	atgggcttgt	2580
gaccctaaac	ttgcaggcaa	ggttagctgc	tttagtaagt	agaattttgt	ggtctttttg	2640
ttttttacat	agtaccaagc	cttgataatt	atgaatttt	tatccattac	taacctttaa	2700
tttaatcaat	catgtacttt	agtttaatgt	ataaagatcc	tctagaaaat	gataatattg	2760
tgtattaaga	cattccttaa	ttaggacaaa	atggctgctg	tatatttact	atatggagtt	2820
ctgagttaaa	taccatcctt	aatactggga	acagaataca	acccatataa	atcagatgca	2880
ggtggtagtc	acatcaccag	agtgatcagt	ataaattttc	ttggtgtatc	cttttccttt	2940
caacacagtg	cagataagag	ttgaatattg	atatcataca	tttagactgc	tgttctgatt	3000
gcatttatct	ttttcctaca	tcatttagaa	ttttatttcc	ctgattcagt	ttttgctgct	3060
gtgaaacagc	tctgatgaac	actaaatatt	aatttcaatt	agctagattg	tacatacttg	3120
cagatttaac	aaaattttag	ggaaattgaa	aaagacatgt	agaatttgtt	gtcttctgct	3180
aagcacgaaa	agttaagata	tctgcttaca	ttgattttgt	agacacatta	agtcaagatt	3240
tggaatttaa	gtcactggca	ggtatctgtg	cattcataga	acttataaag	gtcccaggat	3300
cacttttaag	ggatttttat	tagtttaaag	gtaaataaag	tcagctgaat	ctacatgtct	3360
cttgttttat	ttctctctaa	acttgaaaac	agtaaatctg	cagatactgt	gaggcacaaa	3420
ttatactgtc	aacctactgt	tgctatggtt	atatactccc	acttcataca	ttaccaagag	3480
tcgatcactg	atttaaaatt	tttaatttct	atagttaaga	tttactgcat	aatatagaat	3540

ataaagttaa	gttaacatac	taacatttct	$\mathtt{cctttggagg}$	aagttttaat	ctacttcagg	3600
atgcatatta	ttatcaagat	actttcatat	acaggatagc	ctaattttat	ttgtttaaat	3660
atgcttaata	tgccccagat	tgcaaatgca	tccagtcagt	aatatcactg	tctgtatgtg	3720
gaggacatgt	tcccatggat	catatgtgaa	gatgtcaata	agcttgcatt	aagccacctg	3780
ctttgtaagt	ggattgatta	ataaataact	tatatttc			3818

<211> 1920

<212> DNA

<213> Homo sapiens

60	ggtcacacaa	gaatgacatg	cagggcccac	atctagcagc	catggtecce	caagtgcagg
120	agaaatagag	aaagtccatg	aggttgagag	gcctgtgtgt	aatcagaggg	ggctgactaa
180	accctcatcc	gtctgcaaac	ccctcatcca	acagcaaaca	agatagccag	gaacagagag
240	tgcaaaagct	cccagactcc	tctcatttca	cagtcagccc	caccctcatc	agtctgcaaa
300	cctccacact	ccctgcccat	ttgccactat	tatacagccc	tcttccctgt	tccagaatgg
360	tgtcttcaat	catgcttcca	tcttcaagcc	aacatagatt	gccccgtaag	gggccaattt
420	tgtaggctgc	tgctgtctaa	agagccttca	acaaaagccc	tactcttagc	ggcttcctat
480	gaggatttgt	ctcccaggca	ctaggcaagg	agggagagcc	gttcccctga	tgtggcctga
540	gagaaggaca	gagtgacaca	gtactgggag	caggactggg	attctaggag	ttgggaagtg
600	tggagacctc	tgtgggcgac	gggtcaccca	attcttgata	gccgggagaa	gaaggctgat
660	agagggaggc	cagggaggag	tgttcctcta	gtcttcagaa	ttcaaggagc	ctgaggagtc
720	tatctgtatt	ggacattaat	ttgccccagt	ccccatggcc	tgctcctgtc	ciglaaccac
780	aggggcaaga	agcccagagc	cctcagctgg	ctgctgggcg	gaatatagaa	tccatgcata
840	agtgcaccca	tcactcgtaa	ctctcctaag	accccagaac	gcacaaaatc	gatacagaat
900	gtccctgtaa	acttgcctca	cctgcagaac	tatgtagcat	agtccttttt	gggcctactt
960	tcctgaagaa	atctatttct	cagttgattc	ccctgacctt	gccaaggctt	ctctccatca
1020	cattigicig	cattgtgttt	aacgccagaa	cgacttgcct	caacggccac	gtcaacagag
1080	cttaagcttt	atacatttca	cattcttttc	ggataattcc	atcaattcct	tcaattgttc
1140	ttgccctgac	cagccagcat	ttcagtgatc	aatttacttc	ggcactgagt	gaatgtttct
1200	gtactgctag	caattccact	gcatttgatg	caggctacct	gaaatgccat	acttagagtt
1260	cttaaacaaa	aactatctgt	cagaaacaga	ttgcaagcaa	aaattttcaa	tattgtttaa
1320	gaacaaaatc	atggaaggaa	aggtcagagg	aaactagagc	cattggatga	atagaagatt
1380	tggtttgccc	caggaatata	ttaccaacat	gagcagacca	ggaaggacag	tacagaattt

tttcagggct	ctgcctgcac	ctggcttagc	tccacacact	ttgattccta	gtgtctctga	1440
ttaaagtttc	acatactcag	tagacagaat	agcttgggtt	agtgactctt	cttgaactaa	1500
tcagtgaggc	tatggtatca	ggaatgggga	tgggtaacag	agacacaaac	cctaatggga	1560
agcctctggc	caaacagcca	cctgaatttg	tgctcattgt	atacagtcaa	tcgcttgcta	1620
ctgcacacac	actcttcttc	tcatacaaac	aaaaagaact	ctccttatct	gctatgaacc	1680
cagccatccc	aaggaaaaaag	gcaaatgcac	ttgcctcttc	cccagtgaga	gccacacaca	1740
gcatctccac	cttctgcatg	ctgaagttat	gtcacaggag	gagggcctca	gcacacgatc	1800
aattttictt	ttggtccatt	ggcagcttct	gtctcagaat	ttttgtgtgt	gtatatttgt	1860
agaaaataaa	gaggtacggt	caaggctaca	gccactactg	accattgctg	ccaccatcag	1920

<211> 3433

<212> DNA

<213> Homo sapiens

60	gcggggacaa	gtcaacggct	tcaacggctc	ggtccgctag	acgaccagga	agtcgccatg
120	cggctgggct	tcgcctgact	tttctggacc	gagtcaggcc	gctgccaggc	gtccgttgag
180	ccgggaagag	aacaaggact	attatgaaga	tccactagga	ttgacccagc	gtgcctgaaa
240	gggaattgaa	ctctgcatcg	agcagaacca	gtggttcccc	tgcaggcgcg	gtgcacgaac
300	tgtaaaagct	ttgcagctcc	caggctgtgt	aacaggggca	tececaacea	aacaaggaat
360	tacattggtt	gtggcaattc	agccgtcttt	cccagctccc	agccgtgtct	ccatacccag
420	ctttggaacg	cacctctcat	tcatttctct	aaaatgatca	acactttttg	acatttagta
480	aatttgtggt	ctaacgaatt	gtttcttctt	agacggttag	ttttgcattg	ggagatgact
540	ccttgcttgc	tcctcagagc	atgctggcag	aggtgacagc	ccagaccgag	ttaagaatag
600	ccttcagtcc	tttgaggagc	tttggtggca	gggctcccgc	tcccctgcct	tctcggcacc
660	ctccctcagc	ccggagccca	ctggccaagg	cctttctggg	tgtgggagcc	cccactgcac
720	ctgcacttgc	gggctgtgtg	gcgggaaccg	agagacacga	gtgtggaggg	ttgcagggag
780	gagcagccag	cccgcactca	cttggtgggc	tgggcgtggg	gagttccggg	gggccatctg
840	ctgcggaggg	gggccagtgg	cttagcacct	caatggggga	tggccccggg	ccagccctgc
900	tcgctgggcc	gctcgatttc	ccggcgctgc	tgctggccca	cccccagcag	tgtactgagt
960	tgagcctcct	ccgccatgcc	gacctgcagc	cagggctcgg	tcccacgggg	ttggctgcct
1020	ccctgctcc	cgagcaccac	gcctccccga	tgcggcctga	tgggctcctg	acccaatcca
1080	ggcacaggac	gcgagcgccc	gctgaggaat	ccacgcaagg	gtcccatcga	acggcgccca
1140	cagctgggct	taggtgaagc	caggatccac	agccccagtg	ctccacctgc	tggcaggcag

cctgagtctg	gtggggacgt	ggagagtctt	tatgtctagc	tcagggattg	taaatacacc	1200
aatcagcacc	ctgtgtttag	ctcaaggttt	gtgagtgcac	cagtcaacac	tctgtatcta	1260
gctgctctgg	tgagggcgtg	gagagtcttt	atgtctagct	caaggattgt	aaatacacca	1320
atcagcactc	tgtgtctagc	tcaaggattg	taaatacacc	aatcggcact	ctgtatctag	1380
ctcaaggttt	gtaaacacac	caatcagcac	cccgtgttta	gctcaaggtt	tgtgagtgca	1440
ccagtcgaca	ctctgtatct	agctgccctg	atggggacgt	ggagaacctt	tgtatctagc	1500
tcagggattg	taaacgcacc	aatcagcgcc	ctgacgaaac	aggccactcg	gctctaccaa	1560
tcagcaggat	gtaggtgggg	ccagataaga	gaataaaagc	gggctgcccg	agccagcatt	1620
ggcaacccgc	tcgggtcccc	ttccacactg	tggaagcttt	gttctttcgc	tctttgcaat	1680
aaatcttgct	actgctcact	ctttgggtcc	acgctgcttt	tgtgagctgt	aacactcacc	1740
atgaagatct	gcagcttcac	tcctgagccc	agcgagacca	cgagcccacc	gggaggaacg	1800
aacaactcca	gacacgccac	cttaagagct	gtaacactca	ccgcgagggt	ccaccgcttc	1860
attcttgaag	tcagtgagac	caagaaccca	ccaattccgg	acacaagact	atgagggact	1920
ttattattct	tacttcaaga	ccattattga	agcaccttca	tttttgggag	gactgtggat	1980
gattatgaat	gacaggctta	ctgaatatcc	tcttgtaatt	aatgcagtaa	aacgcttcca	2040
tatttatcca	gagaattctg	gagtccaagg	aagaccaaga	tcaaggcgcc	agcagatttg	2100
gtgtctggtg	aaggctgctc	tccgcttcca	agatggtgcc	ttgatgttgc	atcttcctga	2160
aggagaggaa	cactgtgtcc	tcacatggca	gacagtagga	gagtaatcat	agcctcctgg	2220
tatcgcacat	tcatgggaat	agtgaattta	tttggactag	aaactaagac	ctgctggaat	2280
gtcaccagaa	tagaacctct	taatgaagtt	caaagctgtg	aaggattgcg	agatcctgct	2340
tgcttttatg	ttggtgtaat	ctttatttta	aatggactaa	tgatgggatt	gttcttcata	2400
tatggaacat	acctaagtgg	tactgaactg	ggaggtctta	ttacagtact	gtgcttcttt	2460
ttcaaccatg	gagaggccac	ctgtgtgatg	tggacaccac	ctctccgtga	aagtttttcc	2520
tatcctttcc	ttgtacttca	gatgtatgtt	ttaactttga	ttctcaggac	ctcaagcaat	2580
gatagaaggc	ccttcattgc	actctgtctt	tccaatgttg	cttttatgct	tccctggcaa	2640
tttgctcagt	ttatacttit	tacacagata	gcatcattat	ttcccatgta	tgttgtggga	2700
tacattgaac	caagcaaatt	tcagaagatc	atttatatga	acatgatttc	agttaccctt	2760
agtticatti	tgatgtttgg	aaattcaatg	tacttatctt	cttattattc	ttcatctttg	2820
ttaatgacat	gggcaataat	tctaaagaga	aatgaaattc	aaaaactggg	agtatctaaa	2880
ctcaactgct	ggctaattca	aggtagtgcc	tggtggtgtg	gaacaatcat	tttgaaattt	2940
ctgacatcta	aaatcttagg	cgtttcagac	catatttgcc	tgagtgatct	tatagcagcc	3000
ggaaccttaa	ggtatacaga	ttttgatact	ttaaaataca	cctgttctcc	cgaatttgac	3060
ttcatggaaa	aagcgactct	gctgatatac	acaaagacat	tattgcttcc	agttgttatg	3120
gtgattacat	gttttatctt	taaaaagact	gttggtgata	tttcgcgtgt	tttagctaca	3180
aacgtttatc	taagatgctg	tctttgcagg	tgccatgcct	acaatggcaa	gtgtcaagct	3240
gtctacacct	catcccattg	tgaatcatcc	acattacgaa	gatgcagact	tgaggcctgg	3300

ttgcagcatg	cttgaaatct	gggatgtgga	agacccttcc	aatgcagcta	accctccctt	3360
atgtagcgtc	ctccttgagc	cgagattgtg	ccactgcact	ccagcctggg	cgacaaatca	3420
agaccccgtc	tcc					3433

<211> 2210

<212> DNA

<213> Homo sapiens

60	gcctcgcgat	cgctgtctga	cttctgctcc	ctccgtgtcc	cgctgtgggc	agaaactctg
120	tcccagaaac	gtgttcgggg	tccctcttcg	cctcttcgtc	ctcctcgtcc	gtaccccgcg
180	cgggatcagc	gtcatccagt	aaattgccga	gaccacaggg	cagtggcccg	cgggttctcg
240	gaggcccgga	ccagcaatcg	ctcctggcac	ctgggcgagc	ggtccccttc	gagcctgcag
300	gcgaccctgg	ccctctcaga	agtccccacc	cgacccagga	agggacccag	caccccgcc
360	ctacaccccg	cccctctatc	aaggctgctg	ctccactaaa	gacaggtctc	gcctcgggct
420	aagtaaacgg	tctttttggt	gcacccattg	tctcctaatg	cactcgggaa	agattccttt
480	cagtttagga	gagtgtgtct	agtcattgga	gggttagctc	aattctgatg	tggcgtgaga
540	ccatcctatt	ttcagatggc	cactgcttta	ttggcagtag	cagacctaac	tcgatggttc
600	tctagctcct	atgccgaatt	icagattttt	ttcttcatat	atttgctctg	ccgaatgtta
660	ctgactccgg	gctttggaac	gtttcactaa	aaaagtcgct	tgcgggccct	gcgtctctat
720	cggcggcgtc	gagttccgcg	gccagaaaga	caagatetea	tctttcgtcc	gaaatctcat
780	cgageteect	tegeceegge	tgatcctggg	gcgcgggttc	cgggctggga	ttcaagcgcg
840	aagatggggc	cgagtctctg	ctctgagggg	acctgagcag	gggaagagga	cagggccacg
900	aaaatgtttt	gccttagtta	aggagaagac	ccctgcccca	cccgtcagga	tccccttct
960	cgaactaaga	gtcttgggag	ttcctgatgg	tgggatgagg	tcagagcact	ctcctgctct
1020	tgagagcggt	attttggggc	tcattttcat	accatcgctc	ggcgctcagg	tgtcactcca
1080	cgagccgggg	gtttggagga	gaggggctca	cgcagtcttt	tgaaaaatac	tacaaaggac
1140	ggaaacacaa	tggcaggtaa	tggagtggac	cttggcttct	ttgacgcctt	ttgagggggg
1200	gacagtcaaa	tggaaaattg	ggacctggga	cagaactgca	cccttgcagg	actccgtttc
1260	ccctacagat	aaattgttat	cgtgcattaa	aggtgggttg	atttatgtag	gagtaacaat
1320	tatgtaactt	ctttgcagca	tcaccaccag	gcactgctgt	agggaccagg	aagtgtagga
1380	atcacaacca	caaatatttg	ttcctgttca	taaaaattca	gtatagtgca	tgataacagt
1440	ccctatagga	actigactag	ccactgcttc	ttctccactt	tctttgttcc	gttacaggtt
1500	eccgttcccc	cgccagcccg	ttcagttgat	atgatattgt	attettaaat	aaaaagaatc

acccctgaa	agtagctggg	acaacaggtg	tgcaccacgt	gctgggctca	tttttgtatt	1560
tttagtagag	atggggtttc	gccatattgg	ccaggcagct	ctcgaactcc	agaactcaag	1620
tgatccgcct	gcctgggcct	ccctacacct	ccctgctggg	attacaggca	tgagccaccc	1680
cacttggcca	tcatttgtat	tttgaattta	aatattttgt	agtcatcact	cttcaaagat	1740
tctgggtttt	gctgcatgtg	catgatggga	ttcaggacat	gctactccaa	agtataacat	1800
cttggggtat	tgaatatttt	atgctaaagg	aatatgagta	aaccacagaa	gcaggaaggt	1860
cactgtcacc	ctcccctgc	ccttcctccc	tgaagtccag	tataagactc	ttatgtgaga	1920
ggtgccctct	ctgtacccag	aggaaaggtg	catccttgat	tctgaagaca	cagggacaca	1980
gagaagagcc	tgaacgcaga	ggcctggcta	cctttccccc	actttgttac	tgttagatca	2040
tacttttatt	gacttatcat	acttctccat	gactattcat	tcttgaataa	acctactgtt	2100
ааааааасаа	aaacaaaaac	tgaggtttag	ctgtttccat	gggccttcat	ttcctttgga	2160
aggcgttcct	gtcatgtaaa	atttacatta	aataaatctg	tatgctcttc		2210

〈211〉 3526

<212> DNA

<213> Homo sapiens

60	gtccactccc	cagagtcctt	ccccagcttc	cctgatgggt	catctcaaat	ttcgggtctt
120	aacacagatg	tctgaccatc	ggttggccac	tcagctgagg	agtgtacacc	aagaccaaaa
180	ccagaggagc	atctcttcag	aaactgagga	atcgaggtga	caagctcccc	caaatctgga
240	ggaaacttcc	ggatgggtca	actigacica	gcttggcctt	caagatgaca	aggetteage
300	ctgagcctca	cagcatccag	aggttggcag	ttgggcacag	agtaaagaca	tacacatcga
360	atcctggtcc	tcacttcacc	acaagattac	aaaaccaagg	tcaggaccct	acacaaggca
420	agtgtctaca	aagccagggg	atcagtcaaa	agtgccaaac	ccagaccgtg	tgagtaaggg
480	ctggcctttt	cttccgcatt	tgctgccctc	acttcagcga	tattgatgtg	catcagccat
540	atatggattg	ggctgattcc	ctgagttggt	agccaagacc	cagaggagca	atttacttcc
600	agattcttcc	gaagaatgat	aagttggctt	gggatgctga	cagatgcata	atgigaaiga
660	gaagccacag	aggtgatgca	tgaaggtgac	caagtcgaac	gcccaacagc	agtctttgga
720	aagctcactc	cagcaaacac	atgicitgaa	aaggctgtct	ggctgtggac	tggggctggt
780	gctccctgaa	cgccatcgcc	tcctcccage	cccaggaccc	tagccctgcc	agaagaagac
840	agcaaaagtg	acagagctgg	caagtttaag	atgcagtgaa	acaaaaagga	gaggctggag
900	ggacctggca	tgtgaggaga	ggggctgtct	agagcccagt	gggctccggg	ctgtgaggct

tgtccgccat	ggtccagcct	gtgtggctgc	tttcctggac	tgctgccatc	tgtctgagac	960
cctgactcgg	gaggctcgag	aggatcaatt	gcatctgtgg	acaacggatg	aagaagagga	1020
cttcgatgac	ctcttcttgg	atgacatgcc	tgtgcggacc	ttgttccccg	agagttggct	1080
ctggaggaag	ttcactctgc	caaagagtaa	atcgggcatc	tcccattacc	ccatctctgt	1140
gaaggtgcca	gattccatca	ccacgtggca	gtttgtggtg	gtcagcctca	aggctggaca	1200
aggtctctgt	gtctcggacc	cctttgagct	gacagttatg	aaatcgttct	ttgtggacct	1260
taagttgccc	tcctccgtga	tcaggaatga	gcaggtccag	atccaagcca	tgttgtacaa	1320
tttcagggat	cgccaggcca	aggtccgagt	ggagttcccc	cacaaggaga	cactgtgcag	1380
tgcgtcaaag	ccaggagcac	catcccacca	ggcaggggtt	cagatccagc	aaacgtccta	1440
tagcatcgtc	ctggaacccc	aaggtcagac	ccagacaaaa	ctggtgccaa	gacaggagtt	1500
cttgaacatg	gtacccgaca	cggaggcgga	agtgtttatc	agtgttcaag	gctacactca	1560
gatgetgace	caccggagtt	cagacggcac	ctaccacacc	tccaagggga	acccaggaag	1620
cacttggctc	acaagctatg	tgttccgcgt	ctttgccctg	gcctactcta	tgatgacgac	1680
ccaagtgctt	agcctgtcct	ctctctgtga	catggccaac	tggatcatca	tegacaggea	1740
ggcagaggat	gggcacttcc	tggagaaggg	ccctgtggtc	atgacatcca	tgcaggatgg	1800
ctaccaaggc	tccgaggagg	atgtatccct	cacagctctt	gtcctaatag	ccctgaatga	1860
gggaaaggag	ttgtgcagac	agaagaattt	gatggccagc	atcgagagag	cccgtggctt	1920
cctggagaga	aaacttcccg	acattcagac	aacctttgcc	gtagccatag	cctcctatgc	1980
actggccctt	gccaacagct	cccaagccaa	cgactgcctg	gacagctttg	ccagccctag	2040
tgggtgtggg	atgcttctta	atcaacctca	gtcatggagt	ggggagggg	tgatctcaaa	2100
tccagcaatg	igitaticat	cacttagtgt	ttcctgagca	tccactgtgt	atcacctcct	2160
gagctcagac	cctgtgagac	caagacaaaa	cccactggcc	agtggatgag	cagaatctgg	2220
gctccctgta	caccattgag	gccacagcct	atgggctcat	gcagaagctg	gagctgggcc	2280
ggtacaatga	gacacacgcc	atagccaagt	ggctactaga	gaagcaggag	ctgggaggag	2340
gcttcaggtc	cacccagacc	acggtggtgg	cccttgaagc	tctgacccgc	ttccgcgaag	2400
ctgtcccctt	caagggcatc	caggatetee	acgtccagat	cagageeece	aagacagccc	2460
tgaatgtgaa	ttggtacatt	gatcacagca	atgcctacca	acagcggtca	gcaaagttcc	2520
ttgcccagga	cgacctagag	atcaaagcca	gtggcaacgg	gagaggcacc	atctcgatcc	2580
tgacaatgta	tcacaagtcc	ccagagtccc	gggaggacaa	ctgcaacctg	taccacctga	2640
atgcgactct	ccacagtgcc	ctagaagaaa	ataaaaaggg	aggtgagact	ttteggetee	2700
ggatggaaac	aaggttccag	aacaattgag	aggccacaat	gactatcatg	gaggtctccc	2760
tgctcacggg	cttctacccc	aaccaggatg	acctcaaaca	gctcacgaat	gatgtggaga	2820
ggtacgcctt	tcagtacaaa	accaagacaa	gtaccagcga	cagcactgtt	gtcctctacc	2880
tggaaaagct	ctcccatgag	aagaacacgg	tgctgggctt	tcgggttcac	aggatgctgc	2940
aggcggagtt	cctgcaggcc	gccciggtca	ccatctacga	ctactacgag	ccttgacagt	3000
gcccatccct	gcagaagccc	agtggccaat	tgaggcagga	ggagctccag	acaacagcat	3060

gtgaggcagg	cgtggatttt	gtgtacaaga	caaagctgga	${\tt atctgtggag}$	gtctctgcct	3120
ccaaccctta	cgtctattac	aacacgcagc	tcgaagacat	cattaagagt	ggtacggacc	3180
ctgccacacc	cctggccatg	aagaaattcg	tctcccatgc	cacttgccat	gactccctgg	3240
ggttgcaaga	acaggaatcg	tacctcatca	tgggccagac	gtcagacctg	tggagaatca	3300
aatctgatta	cagctatgtt	ctgggcaagg	agacgttcct	catcctttgg	ccagcagatg	3360
gagatgccag	caagaaagaa	ttgcgggacc	aactggagga	atttttggaa	tatatgcgca	3420
cccacggctg	ccagtcctga	gcctcttctg	ctttcaggga	ggtgtcatca	ggcagctctg	3480
ggccactggg	tttaacccca	aataaagagc	acaggatatg	acaccc		3526

<211> 3315

<212> DNA

<213> Homo sapiens

gcttccgggt	gagccccccc	cgctcttacg	cggtctgtgg	gagaccggag	cgggagacag	60
cggtgacagg	agcagcggcc	gggagccctt	agggaggcag	acagagcctg	cagccaatgc	120
cccaggagcc	ctcggttcca	accaactgat	gccctgtgc	ccactggccc	acgccatgca	180
gccccagtcc	gttctgcaca	gcggctactt	ccacccacta	cttcgggcct	ggcagacagc	240
caccaccacc	ctcaatgcct	ccaacctcat	ctaccccatc	tttgtcacgc	cctgggcttg	300
gccaggcagg	gaagccagac	actggatccc	atcctcctcc	caccatctcc	acttccatat	360
ttctttcctg	cttcccaacc	atccctctca	gtcgcccccg	caccactggc	ccttcccaca	420
gctaccaatc	catateccae	ccccgctctt	gcagggatgt	tcctgatgac	atacagccta	480
tcaccagect	cccaggagtg	gccaggtatg	gtgtgaagcg	gctggaagag	atgctgaggc	540
ccttggtgga	agagggccta	cgctgtgtct	tgatctttgg	cgtccccagc	agagttccca	600
aggacgagcg	gggttccgca	gctgactccg	aggagtcccc	agctattgag	gcaatccatc	660
tgttgaggaa	gaccttcccc	aacctcctgg	tggcctgtga	tgtctgcctg	tgtccctaca	720
cctcccatgg	teactgeggg	ctcctgagtg	aaaacggagc	attccgggct	gaggagagcc	780
gccagcggct	ggctgaggtg	gcattggcgt	atgccaaggc	aggatgtcag	gtggtagccc	840
cgtcggacat	gatggatgga	cgcgtggaag	ccatcaaaga	ggccctgatg	gcacatggac	900
ttggcaacag	ggtatcggtg	atgagctaca	gtgccaaatt	tgcttcctgt	ttctatggcc	960
ctttccggga	tgcagctaag	tcaagcccag	cttttgggga	ccgccgctgc	taccagctgc	1020
cccctggage	acgaggcctg	gctctccgag	ctgtggaccg	ggatgtacgg	gaaggagctg	1080
acatgeteat	ggtgaagccg	ggaatgctct	acctggacat	cgtgcgggag	gtaaaggata	1140
agcaccctga	cctccctctc	gccgtgtacc	acgicicitgg	agagtttgcc	atgctgtggc	1200

atggagccca	ggccggggca	tttgatctca	aggctgccgt	actggaggcc	atgactgcct	1260
tccgcagagc	aggtgctgac	atcatcatca	cctactacac	accgcagctg	ctgcagtggc	1320
tgaaggagga	atgatggaga	cagtgccagg	cccaagaact	agaactttaa	aacgttcccg	1380
gggcctcaga	caagtgaaaa	ccaaagtaaa	tgctgctttt	agaactgtgc	cctcatgccc	1440
tcttcctgct	cacatgctag	cggggcccag	cagccctggg	tggttttgcc	agcatgctaa	1500
ctcttgtaac	tcgcagctgc	atcctatgag	ctctcccaag	cttccccgcc	cctccctgg	1560
gtcagccgtg	aggcccacct	ttgccaccct	cagctctttc	ctctggtgtg	gcttcagctt	1620
gaaagcaacc	tggagtcggg	ggcacagcct	ttggggcctg	gctgggagag	ggtcttggag	1680
cattagggga	agaagagagc	agtgggatct	tggggcctga	gaagccttgg	aacgcttctg	1740
gcagcagagc	tgggtgtggg	aatgaggcct	agatcgatat	ccctgggtta	gagttgaaat	1800
ttgccgcaat	tccactggaa	ggcatttccc	acgaggccag	aggttgccag	gctgcctgag	1860
gtctcctatt	ctactctgaa	ccataaaccc	agagaagaat	tactcattaa	ccagcataaa	1920
tactgcctga	ggatcaaaac	tcagaggcaa	agagggagtt	cctgactgct	agaggtgcca	1980
ccaccacaaa	cactttttat	tcaggagata	ctttttgaga	atctctgctc	tgttcctagg	2040
ttcagtgctg	ggtcctggga	atacagcagg	acagacctca	gcttatctct	tcatagaaat	2100
tatacaaaga	gaattgggga	gacagctaag	aagaaaacaa	agaaataaag	cagttacaaa	2160
ttgtgataag	tgctttgaag	gaaagaaggg	gtctgagaca	acaacaggga	aggggcctct	2220
cttgaaacag	tagttgggaa	ggaggcagac	atgcaccagt	gatgtggtga	caggtgctct	2280
gaaggaggtc	accaggacct	gacctctttg	aaggatcaga	aaatacttcc	ctgaaggact	2340
gacatttgag	cctagacctg	aagggtgagc	catcaagcta	agacaattgg	ggaagagcat	2400
tccagggaga	gggaggagtt	gtgcaaaggc	cctggggctc	cttctagctg	gaggaatgca	2460
aggctagctt	gtctggagca	ctgagaggat	ggcctgaact	gagtggagag	agacagacca	2520
ggaccaaacc	atgcagaggt	caagggccac	attcaccttt	tcagagtgac	tcaatcaaat	2580
ttgtagtttg	taaaagtatt	ttaacagctc	tgcggcaaag	tgcaaatgaa	aagtettgat	2640
ggcatggact	ggagcgggga	cagtggggat	ggagaaaggg	gaatggattg	tggatgtgtt	2700
tagaaggtag	attcgatgtg	aaggatgaat	ctggcttgac	cttctgggtg	gctgatgggc	2760
catttactga	gatggggcag	cctggaagag	gaacagaagc	agggtcgggg	tggagggaga	2820
atactaaact	tagcttgaga	cattttgcaa	taaggaagct	atatctagag	tgcttatgtg	2880
actcacctaa	ggccactcaa	caagtttgtg	gcagaactgg	attagaactg	cacagaaaac	2940
agccaagctg	ggatttgaac	ccatglagic	caactccaag	gcctctgccc	ctaaccactg	3000
tgccatacca	cctcccaata	atcaacagca	aaattatagg	tctaacaatg	ttttatagac	3060
acccctccat	ttatgtgatg	ggtttgcatc	ctgataaacc	catcataagt	tgaaaatatg	3120
atcataagtt	gaaaatatga	tcataagtca	aaaatgtatt	taatatacct	aacctaccaa	3180
acateatage	ttagcctagc	ctgccttaaa	catgctcaga	acacttacat	tagcctacag	3240
tgggcaaaac	tatccaacac	aaaatctata	tigiaataaa	gttataaaga	attttgaata	3300
aaaattcaat	atttg			•		3315

```
<210> 4653
<211> 2322
<212> DNA
<213> Homo sapiens
```

60	acaaagtggt	gggagactga	ccagcagatg	tatttcagat	ggccttctgt	aagtggcctt
120	tcactggctt	tttgaaccgt	cccactctgg	tctacactca	caccagtttt	atcctgggcc
180	gaggtgttct	ccacaggggt	ctgtgtgtag	tatgtcaaag	tgagcccaac	ccagaacatt
240	aggaggcctg	tgatgtgggg	gctcaggcta	agaacaaagg	gctcttcctg	tgggagagct
300	gtgcctcctc	gcttatcaag	ttcccaccct	gcagctgaag	ctatgcctgg	gacctgtgtc
360	acctcaagcc	cacttgcaga	ccaacttctt	tttgacccaa	ctagtctcta	tgggaacccc
420	catggagaag	aggatgataa	ttgtttattc	ctccagtgga	tgacttcttc	ctaatatgta
480	gccctttcag	tctggattgg	gccttccctt	ataccctcgt	ttattgaaaa	cttgaggaaa
540	aacagatccc	ttctgagcag	gcaaagacac	cccagactat	gtatctatga	gcatttttct
600	agcggctcta	gaaaaggact	ccacttcttg	attctcacct	acctgcagaa	aagtcccagt
660	ttttaacatc	ctggattcca	ctactaactc	gcatcgtcgc	agtggttcca	gacggaccca
720	taagtgggag	tgatgctgga	tctgtgaaaa	gatggctcat	acattgaggt	ctgaaagcat
780	ctcgatgtct	agcacatcaa	gaggtctatg	cacaagcgtg	gcactcagga	aagatttgca
840	aaacagcacc	actgccagac	aaggagacca	cgctttcagc	tcatgaaatg	ctggatatag
900	ccgcttgtac	tcatatttca	ctcagcaaaa	catatttgaa	atgcaaaagc	catgatectt
960	ccgcttccag	ctcagggcta	aaactcagcc	cataattttc	atcacagtga	agtttgttgt
1020	aaagaaatcc	tccaggaaag	gatacaataa	tcagtacaca	gagtgttgaa	aagttaagcc
1080	ttttctggat	agtaccagga	ccgaagagga	ggataacact	gggtaaagca	ctccaggctg
1140	tgtacactct	cagatattga	agcagcttct	tgaaagtggt	ctgccaagga	attgtccttt
1200	ctcctggatc	cagcaagcat	gacaccttgg	ggcaggacat	cattcctgtt	gaagtgagca
1260	ggtcaggggc	gccgggagga	caagagagat	ccctgagcat	tggctctgaa	ctttactgcc
1320	gtacaccaca	gtgagatgtc	gaccagctgg	tatcactigg	atgggtcttc	atcctggggg
1380	cagagatete	cgtccatttc	cctgcagtcc	ccgattgatt	aggagacgtg	atgtgcatca
1440	cgtggttctt	cagggatcac	acattgcctg	agatggatgc	ttaccttccc	agcaagccac
1500	ctttgacccc	acccaaaggt	gtctggaaaa	caaccctgct	gtcttcacca	agtatttggg
1560	accattctca	atgectactt	agacacccct	ttctgatcag	ctcaggagaa	ttgaggttct
1620	ggtaaccatt	ttgagttaaa	tttgccatga	tgggcaggag	ggaactgcat	gctggatcaa
1680	tactttcccc	ccaggcctct	ccagacccca	cagagtgact	tgctccactt	gccttgattc

aaccatttta	tcctcaagcc	caagaatggg	atgtatttgc	acctgaagaa	actctctgaa	1740
tgttagatct	cagggtacaa	tgattaaacg	tactttgttt	ttcgaagtta	aatttacagc	1800
taatgatcca	agcagataga	aagggatcaa	tgtatggtgg	gaggattgga	ggttggtggg	1860
ataggggtct	ctgtgaagag	atccaaaatc	atttctaggt	acacagtgtg	tcagctagat	1920
ctgtttctat	ataactttgg	gagattttca	gatctttct	gttaaacttt	cactactatt	1980
aatgctgtat	acaccaatag	actttcatat	attttctgtt	gttttaaaa	tagttttcag	2040
aattatgcaa	gtaataagtg	catgtatgct	cactgtcaaa	aattcccaac	actagaaaat	2100
catgtagaat	aaaaatttta	aatctcactt	cacttagccg	acattccatg	ccctgaccaa	2160
tcctactgct	tttcctaaaa	acagaataat	ttggtgtgca	ttctttcaga	ctttttccta	2220
tacattttat	atgtagaaat	gtagcaatgt	atttgtatag	atgtgatcat	tcctatattg	2280
ttattgattt	ttttcactta	ataaaaattc	accttattcc	tt		2322

<211> 4478

<212> DNA

<213> Homo sapiens

gttgcacttt	gcaaaggagg	agtcagccct	catgtgcaag	aagctcacta	agcttgccaa	60
ggagaatgac	agcatgaagg	aggagctgct	gaagtaccgc	tegetetatg	gggacctgga	120
cagcgcgctg	tcagccgagg	agctggccga	tgcccccac	tegegggaga	ccgagctgaa	180
ggtgcacctg	aagctggtgg	aggaggaagc	caacctgctg	agccgccgca	tcgtggagct	240
ggaggtggag	aaccgaggcc	tgcgggctga	gatggacgac	atgaaggatc	atggaggtgg	300
ctgtgggggt	cctgaggcac	gcctggcctt	ciccgcgctg	ggtggcggag	agtgcgggga	360
gagcttggca	gagctgcggc	gacacctgca	gtttgtcgaa	gaggaggccg	agctgctgcg	420
gcgctcctct	gccgagctcg	aggaccagaa	caagcigcig	ctgaacgagc	tggccaagtt	480
ccgctcggag	cacgagctgg	acgtggcgct	gtcggaggac	agtigitcig	tgctcagcga	540
accttcacag	gaggagctgg	cggccgccaa	gctgcagatc	ggcgagctca	gcggcaaggt	600
caagaagctg	cagtacgaga	accgcgtgct	cototocaac	ctccagcgct	gtgacctcgc	660
ctcctgccag	agtacgcggc	ccatgctgga	gacggacgcc	gaggccgggg	actctgccca	720
gtgtgtgcct	gctccctgg	gegagacaca	cgagtcccat	gcggtccgac	tctgcagagc	780
cagggaggcc	gaggtgctgc	ctgggctgag	agagcaggcc	gccctggtca	gtaaggccat	840
cgatgtcctg	gtggctgatg	ccaatggct1	caeggetgge	ctccggctgt	gtctggacaa	900
cgagtgtgct	gacttccggc	tgcatgaggc	ccccgacaac	agcgagggcc	ccagggacac	960
caageteate	catgccatcc	tggtgcgcct	gagcgtgctg	cagcaggagc	tgaatgcctt	1020

cacgcggaag	gcagatgcag	tcctcgggtg	ctctgtcaag	gaacagcagg	agtccttctc	1080
atcactgccc	cccttgggct	cccaggggct	ctctaaggag	attcttctgg	caaaagacct	1140
tggctcagac	tttcagccac	ctgacttcag	ggacctgccg	gaatgggagc	ccaggatccg	1200
agaggctttc	cgcactggtg	acttggactc	taagcccgac	cccagccgga	gcttcaggcc	1260
ttaccgagct	gaagacaatg	attcctatgc	ctctgagatc	aaggagctgc	agctggtgct	1320
ggctgaggcc	cacgacagcc	tccggggctt	gcaagagcag	ctctcccagg	agcggcagct	1380
acgaaaggag	gaggccgaca	atttcaacca	gaaaatggtc	cagctgaagg	aggaccagca	1440
gagggcgctc	ctgaggcggg	agtttgagct	gcagagtctg	agcctccagc	ggaggctgga	1500
gcagaaattc	tggagccagg	agaagaacat	gctggtgcag	gagtcccagc	aattcaagca	1560
caacttcctg	ctgctcttca	tgaagctcag	gtggttcctc	aagcgctggc	ggcagggcaa	1620
ggttttgccc	agcgaagggg	atgacttcct	cgaggtgaac	agcatgaagg	tgctgtactt	1680
gctgatggag	gaagaggaga	taaacgctca	gcattctgat	aacaaggcct	gcacggggga	1740
cagctggacc	cagaacacgc	ccaatgagta	catcaagaca	ctggccgaca	tgaaggtgac	1800
gctgaaggag	ctgtgctggc	tgctccggga	tgaacgccgt	ggtctgacgg	agcttcagca	1860
acagtttgcc	aaggccaagg	ctacctggga	gacagagcgg	gcagagctca	agggccatac	1920
ctcccagatg	gagctgaaga	cagggaaggg	ggccggggag	cgggcagggc	ccgactggaa	1980
ggcagcccta	cagcgggagc	gtgaggagca	gcagcacctc	ctagctgagt	cctacagcgc	2040
tgtcatggag	ctgactcggc	agctgcagat	cagtgagcgc	aactggagcc	aggaaaagct	2100
gcagctggtg	gagcggctgc	agggtgagaa	gcagcaggtg	gagcagcagg	tgaaggagct	2160
gcagaaccgc	ctaagccagc	tgcagaaggc	tgccgacccc	tgggtcctga	agcactcgga	2220
gctggagaag	caggacaaca	gctggaagga	gacacgcagt	gagaagatcc	acgacaagga	2280
ggctgtttcc	gaagttgagc	ttggaggaaa	tggtttaaag	agaaccaaat	ctgtttcttc	2340
catgtctgag	tttgaaagtt	tgctcgactg	ttccccttac	cttgctggcg	gagatgcccg	2400
gggcaagaag	ctgcctaaca	accctgcctt	tggctttgtg	ageteegage	caggggatcc	2460
agagaaagac	accaaggaga	agcctgggct	ctcgtcgagg	gactgcaacc	accigggige	2520
cctggcctgc	caggaccccc	cagggaggca	gatgcagcgc	agctacacgg	ctcctgacaa	2580
gacgggcatc	cgagtctact	atagtccccc	ggtggcccgg	cgcctcggag	tccctgtggt	2640
tcatgacaaa	gagggcaaga	tcattatcga	gcccggcttc	ctcttcacca	cagccaagcc	2700
caaagagtcg	gccgaggctg	atgggctggc	tgagagetee	tatggtcggt	ggctctgcaa	2760
cttctcacgg	cagcgcctgg	acggaggctc	agcgggcagc	ccctcggcgg	ccgggcctgg	2820
cttcccagcg	gccctgcatg	actttgagat	gtcaggcaac	atgagtgatg	acatgaagga	2880
gatcaccaac	tgtgtgcgcc	aggccatgcg	ctccggctca	ctggagagga	aagt gaagag	2940
cacatccagc	cagacggtgg	gcctggccag	tgtgggcaca	cagaccatcc	gcacggtcag	3000
cgtgggcctg	cagaccgacc	caccccgcag	cagectecat	ggcaaggcct	ggtcaccccg	3060
cagctcttcg	ctcgtgtctg	tgcgcagcaa	gcagatetee	tectecetgg	acaaggtcca	3120
ttcgcgcatc	gagcggccct	gctgctcccc	caagtatgge	tcaccaaagc	tccagaggcg	3180

```
gtctgtgtcc aagctggaca gcagcaagga ccgcagcctg tggaacctgc accagggcaa
                                                                 3240
                                                                 3300
gcagaacggc tcggcctggg cccgctccac caccacgcgg gacagccctg tattgagaaa
                                                                 3360
catcaacgat ggacteteca geetetteag tgtggtggag cactcaggga geacggagte
tgtctggaaa ctaggcatgt ctgagacgcg ggccaagccc gagcctccca agtacggcat
                                                                 3420
tgtgcaggaa ttcttccgta atgcgtgtgg ccgggcaccg agccccacct catcagcagg
                                                                 3480
agaggagggc accaagaagc cagagcccct ctccccagcc agctaccatc agccagaggg
                                                                 3540
tgtggccagg atcctgaaca agaaggcagc caagttgggc agcagtgagg aggtcagact
                                                                 3600
caccatgctc ccccaggtgg ggaaggatgg tgtcctccgg gacggagatg gagccgtggt
                                                                 3660
cetteceaat gaggaegetg tttgtgaetg tagtaeceag teteteaeet eetgettege
                                                                 3720
ccgatcgtcc cgctctgcca tccgccactc tccttccaag tgcaggctgc acccttcaga
                                                                 3780
gtccagctgg ggtggggagg agagggcact ccccccagc gagtgacaga gcagccaagc
                                                                 3840
3900
                                                                 3960
ggctgaacac cagggtcatg ccagtcccgc cagccgcctc ctccatgccc cagtgactgg
                                                                 4020
tgtgggcaga gcaggcagcc agtggagcig tgggccagii ccgcictigg atgcigctgc
                                                                 4080
teteacecat gaggteaggg gggeeeteea aggttatete eaggtgaggg gatteacate
aggccacaag ccaccagagg ccttctgcca cctcccagag cgacagccag ggaggctgcg
                                                                 4140
tactcagcct cggggagaaa tccccgtggg acctgagccc caagacctac ggaccactca
                                                                 4200
gccttaccat cgtaccgtcc aggattgtcc ttgccatctt tgttgtctca gccagacctt
                                                                 4260
                                                                 4320
ggttttcagt aaagccccag tttctacttc ctgcatgcca ctgtgcaagg ccactcatca
                                                                 4380
ctgttcctgc agaagcetct ggacgtgggg ctggatgggg ttgaaaatgt tacatgtaaa
tattggtttg gttcggtttt tagcatttta cttggtaact ggttgttttc ttttttgggg
                                                                4440
                                                                 4478
tggggggatt ggtttgtaaa aattctctac tcttttgg
```

<211> 2854

<212> DNA

<213> Homo sapiens

agaactcaca	tcggatgatt	caggcatggc	tctgctaaca	ctttattaaa	agcatggatt	60
aattttactt	ccaagtttat	tittacigca	ccgtcccatt	tgtggaaaca	actagettae	120
tcagcttttt	tttcctttta	taaaggaaag	aacagaaaag	taaaaggagg	aaagaaaaca	180
agaggtgagt	gaggcaactg	aaaactgttc	ttggacctgc	ggtgctatag	agcaggctct	240
tctaggttgg	cagttgccat	ggaatctgga	cccaaaatgt	tggcccccgt	ttgcctggtg	300
gaaaataaca	atgagcagct	attggtgaac	cagcaagcta	tacagattct	tgaaaagatt	360

tctcagccag	tggtggtggt	ggccattgta	ggactgtacc	gtacagggaa	atcctacttg	420
atgaaccatc	tggcaggaca	gaatcatggc	ttccctctgg	gctccacggt	gcagtctgaa	480
accaagggca	tctggatgtg	gtgcgtgccc	cacccatcca	agccaaacca	caccctggtc	540
cttctggaca	ccgaaggtct	gggcgatgtg	gaaaagggtg	accctaagaa	tgactcctgg	600
atctttgccc	tggctgtgct	cctgtgcagc	acctttgtct	acaacagcat	gagcaccatc	660
aaccaccagg	ccctggagca	gctgcattat	gtgacggagc	tcacagaact	aattaaggca	720
aagtcctccc	caaggcctga	tggagtagaa	gattccacag	agtttgtgag	tttcttccca	780
gactttcttt	ggacagtacg	ggatttcact	ctggagctga	agttgaacgg	tcaccctatc	840
acagaagatg	aatacctgga	gaatgccttg	aagctgattc	aaggcaataa	tcccagagtt	900
caaacatcca	attttcccag	ggagtgcatc	aggcgtttct	ttccaaaacg	gaagtgtttc	960
gtctttgacc	ggccaacaaa	tgacaaagac	cttctagcca	atattgagaa	ggtgtcagaa	1020
aagcaactgg	atcccaaatt	ccaggaacaa	acaaacattt	tctgttctta	catcttcact	1080
catgcaagaa	ccaagaccct	cagggaggga	atcacagica	ctgggaatcg	tctgggaact	1140
ctggcagtga	cttatgtaga	ggccatcaac	agtggagcag	tgccttgtct	ggagaatgca	1200
gtgataactc	tggcccagcg	tgagaactca	gcggccgtgc	agagggcagc	tgactactac	1260
agccagcaga	tggcccagcg	agtgaagctc	cccacagaca	cgctccagga	gctgctggac	1320
atgcatgcgg	cctgtgagag	ggaagccatt	gcaatcttca	tggagcactc	cttcaaggat	1380
gaaaatcagg	aattccagaa	gaagttcatg	gaaaccacaa	tgaataagaa	gggggatttc	1440
ttgctgcaga	atgaagagtc	atctgttcaa	tactgccagg	ctaaactcaa	tgagctctca	1500
aagggactaa	tggaaagtat	ctcagcagga	agtttctctg	ttcctggagg	gcacaagctc	1560
tacatggaaa	caaaggaaag	gattgaacag	gactattggc	aagttcccag	gaaaggagta	1620
aaggcaaaag	aggtcttcca	gaggttcctg	gagtcacaga	tggtgataga	ggaatccatc	1680
ttgcagtcag	ataaagccct	cactgataga	gagaaggcag	tagcagtgga	tcgggccaag	1740
aaggaggcag	ctgagaagga	acaggaactt	ttaaaacaga	aattacagga	gcagcagcaa	1800
cagatggagg	ctcaagataa	gagtcgcaag	gaaaacatag	cccaactgaa	ggagaagctg	1860
cagatggaga	gagaacacct	actgagagag	cagattatga	tgttggagca	cacgcagaag	1920
gtccaaaatg	attggcttca	tgaaggattt	aagaagaagt	atgaggagat	gaatgcagag	1980
ataagtcaat	ttaaacgtat	gattgatact	acaaaaaatg	atgatactcc	ctggattgca	2040
cgaaccttgg	acaaccttgc	cgatgagcta	actgcaatat	tgtctgctcc	tgctaaatta	2100
attggtcatg	gtgtcaaagg	tgtgagctca	ctctttaaaa	agcataagct	ccccttttaa	2160
ggatattata	gattgtacat	atatgctttg	gactattttt	gatctgtatg	tttttcattt	2220
tcattcagca	agttttttt	tttttcagag	tcttactctg	ttgcccaggc	tggagtacag	2280
tggtgcaatc	tcagctcact	gcaacctctg	cctcctgggt	tcaagagatt	cacctgcctc	2340
agccccctag	tagctgggat	tataggtgta	caccaccaca	cccagctaat	ttttgtattt	2400
ttagtagaga	tggggtttca	ctatgttggc	caggctggtc	tcgaactctt	gacctcaaat	2460

gatccacccg	cctcggcctc	ccaaagtgct	gggtttacag	gcatgagcca	ccatgcccag	2520
ccctcattta	gcaaagtttt	aaacatgaaa	agtgcttatt	agaggatatc	agtgcctggc	2580
ccacatgaga	gaacagatcc	atacacactt	tgaaaaactt	tgttcacttt	taggaaatat	2640
aattttgaaa	aatcatttac	atacaagagg	tccactgagg	cattgctttt	aatggcaaaa	2700
tattgcaatg	tacttgaatg	tccttcacat	tagattggta	agataaattt	tagtatgtgc	2760
atgtactgga	atattatata	gccagtaaac	aaattgacaa	tgaagctcta	tttgtaccag	2820
taaagaatgg	tcttgaagag	acattgtaaa	atga			2854

<211> 3284

<212> DNA

<213> Homo sapiens

attaagctga	gaccagaagg	ctgagtggtg	gtgacctaca	agaagaagaa	gggaaggtca	60
taacagtaat	gacttggagt	ggagaggccc	tgcaggttgg	agaagttcag	tgtggctcac	120
catggagtgt	cagagtgagg	gagatgactg	ctgagctccg	agaggtgaac	agagcccctg	180
aattaattaa	aaggatttac	ggtatatctt	atgactctac	aggagccttg	ggatatagat	240
aaaaatgaat	gccatttccc	atccagtact	gaagtgttaa	agttagaaga	catttgctaa	300
gtcaaatcac	ttcaggtatg	ttgaagaaag	tacgcacatc	ttatcttctg	aacattttga	360
tggaattaac	cacaaacttc	atgcaagaac	agtaaaagga	aagatactat	tcaagtccag	420
tacatcataa	tgcagaattc	tgtatcagta	ccaccaaaag	atgaagcaag	caaggagaag	480
cttaggagaa	aaaataatgg	aacaagtatc	aggaaagctg	gactccattc	catacctgcc	540
tccaaccagc	tgtggaaccc	caggcaaagt	tattttgccc	ctgagcctca	gttttctcat	600
tgaaaaaatcg	agaatgacta	ggtgttcctt	tcaggctcta	aaaaggtgaa	tcaaacatcc	660
cttctggaac	aatacagagt	aggaaaggtt	tgcagaataa	gagtcagttt	aggaccattg	720
caccaaaaaat	tgtgcccaaa	gtcctaacgt	ccagaatgct	gccatgtcat	tcaccatcac	780
gctctgatca	ggtgaatctg	ggaccctcca	tcaactccaa	gctgctgggg	atgtccaccc	840
agaactatgc	cctgatgcag	gttgctggcc	aggaggggac	attttctctt	gttgctctgc	900
cacatgttgc	ctcagctcag	ccaattcaga	aacccagaat	gtccctacct	gaaaacctga	960
aacttcctat	tcctagatat	caacccccta	gaaatagcaa	agcatcaaga	aagaaaccca	1020
tcctgatctt	tcctaagagt	ggctgtagca	aagctcctgc	ccaaacccaa	atgtgtcctc	1080
agatgtcccc	ttccccacct	caccaccctg	aactcctgta	caaacccagt	ccatttgaag	1140
aagtaccatc	actagagcaa	gccccagcca	gcattagcac	agctgcgctg	accaatggaa	1200
gtgaccatgg	ggacttgaga	ccaccagtga	ccaacaccca	tggcagtctg	aaccctcctg	1260

ctaccccagc	atcatccaca	ccagaggagc	ctgccaagca	ggacctcaca	gctctttcag	1320
ggaaagcaca	ctttgtaagc	aagataacat	ctagtaaacc	ttctgctgtt	gccagtgaaa	1380
aatttaaaga	acaagttgat	cttgcaaaaa	ccatgaccaa	tttatcacca	accattcttg	1440
gcaatgcagt	tcagttgatc	tcttcagtcc	ccaaagggaa	actgccaatc	ccaccctact	1500
caagaatgaa	gacaatggag	gtttacaaaa	tcaaatcaga	tgctaacatt	gcaggttttt	1560
ctttaccagg	acctaaggcc	gactgtgata	agataccctc	caccacagaa	ggctttaatg	1620
cagccaccaa	ggtggcaagc	aggctacctg	ttccacaagt	gtcacagcag	agtgcctgtg	1680
aaagtgcctt	ttgtccaccc	accaaacttg	atcttaacca	caaaacaaag	ctgaacagtg	1740
gagcagcaaa	gagaaaagga	agaaaacgga	aggtaccaga	tgaaattttg	gcatttcagg	1800
gaaaaaggag	gaaatatatc	attaataagt	gtagagatgg	taaagaaaga	gtaaaaaaatg	1860
atccccaaga	attcagagac	caaaagctgg	ggaccctgaa	aaaataccgt	agcattatgc	1920
ccaaacctat	catggtcata	cccactttgg	cctccctggc	ttctccaact	acactacagt	1980
cccagatgct	tgggggccta	ggacaggatg	ttttgttaaa	taattcactc	actcctaaat	2040
atcttggctg	taagcaagac	aacagctctt	cccctaagcc	cagctccgtg	ttcagaaatg	2100
gattctctgg	cattaagaag	ccttggcaca	gatgtcacgt	ctgcaaccac	cacttccagt	2160
tcaaacagca	ccttcgagac	cacatgaata	cacacaccaa	cagacgccct	tacagttgtc	2220
ggatttgtcg	caagtcctat	gtacgtcctg	gcagcctgag	cacacacatg	aaacttcatc	2280
atggtgagaa	ccgtctgaag	aaactcatgt	gttgtgagtt	ttgtgcaaaa	gtgtttggcc	2340
acatccgagt	ctattttggc	catctgaaag	aagtgcatag	ggttgtgatc	agcactgagc	2400
ctgcgcccag	tgaactgcag	ccaggagaca	taccaaagaa	cagagacatg	agtgtgcgag	2460
gcatggaggg	atcattggag	agggaaaaaca	agtccaacct	ggaagaagac	ttccttctaa	2520
accaggcaga	cgaagtcaaa	ttacaaatca	aatgtggtcg	ttgtcagatt	actgctcagt	2580
ctitigegga	aataaaattt	catttacttg	gtgttcatgg	agaggaaatt	gagggcaggc	2640
tacaagaagg	gaccttccca	ggaagcaagg	ggactcagga	agagttggtg	cagcacgcta	2700
gccccgactg	gaaaaggcat	cctgagagag	ggaagccgga	gaaggttcat	tcctcctccg	2760
aggaatcaca	tgcatgtcca	agactgaaaa	ggcagctcca	ccttcatcag	aatggcgtgg	2820
aaatgctcat	ggaaaatgaa	ggaccccagt	caggaaccaa	caagccaagg	gaaacctgcc	2880
agggccctga	gtgtcctggc	ctccacacgt	ttctcttgtg	gtcccattca	ggctttaact	2940
gcctgctttg	tgcagagatg	ctgggacgga	aagaggacct	cctccaccac	tggaagcacc	3000
agcataactg	tgaggaccct	tccaaactgt	gggctatttt	aaatacggtc	tccaaccagg	3060
gagtgatcga	actttccagt	gaagctgaga	aatgagaccc	caaggcagcc	tggggttaag	3120
gagagagctc	tgccgccacc	ttccttcaga	gcttcgtgct	ttatggtggt	gcttagtcac	3180
aaagatcaaa	caacaggatt	ggtgtgagtg	aacagaaatg	atttttgtac	atggttttat	3240
tttcttaacg	aaataaaata	taagctctcg	aagcatattt	ttct		3284

<210> 4657 <211> 2474 <212> DNA <213> Homo sapiens

<400> 4657

60 ttaaaaacta ggcttggcct cttgcctata gtggccacca ctcctcaagc cccagccagc 120 accatgageg geegagttgg agacetgage eccaaacagg cagagacect ggeeaaggtt caagcaattc tcctgcctca gcctcctgag tagctgggat tacagttccg agaaaacgtc 180 240 caggatgtgc ttcctgccct gcccaaccct gatgattatt tccttctacg ctggctccga 300 gctcggaatt ttgacttgca gaagtcggag gctttgctcc gcaagtacat ggagttccgg 360 aagaccatgg atattgacca tacccttgat tggcagcccc cagaggtgat ccagaagtac 420 atgcctgggg gcctgtgtgg ctatgaccgt gatggctgcc ccgtgtggta tgacatcact 480 gggccacttg atcccaaggg gtlgctcttc tcagtcacca agcaggacct gctcaagacc 540 aagatgaggg actgtgagcg catcctgcat gagtgtgacc tgcagacaga gaggctaggg aagaagattg agaccatcgt gatgatattt gactgtgagg gcctgggact gaaacacttc 600 tggaaacctc tggtagaagt gtaccaggag ttctttggcc tccttgaaga gaattaccca 660 720 gagaccctga agttcatgct catcgtgaaa gctaccaaac tgttccctgt gggctacaac 780 ctcatgaagc cattcctgag tgaggacact cgcaggaaaa ttattgtgtt gggaaataac tggaaggaag gtttgctgaa acteateagt eetgaggaac tgeetgeeca gtttggggge 840 900 accetgactg acceagatgg gaaccecaaa tgtttaacca agattaacta tggcgggag atecceaagt ceatgtacgt gegggaccag gtgaagacte agtacgagca eteggtgeag 960 1020 atcaaccgcg gclcatcaca ccaagtggaa tacgagatcc tattlccagg ctgcgttctc 1080 aggtggcagt tctcatctga cggtgcggac atcggcttcg gagttttcct gaagaccaag 1140 atgggggage gacageggge aggggagatg acagaggtte tacccageca gegetataac 1200 gcccacatgg tgcccgagga tgggaacctc acctgctcag aggccggcgt ctatgtggaa 1260 agigagicit ggaagicit cigicalci cccgigatta taigcagcca tgagciacag 1320 aactcacact caaactcgca agtgatggca taccaaatgg tacgaaaatg caagctaagt 1380 aggeetttae eceteeetge eagtaactaa tittigteig igigagtaac iggeetgigt 1440 gtaagaggac teacceceta egiggietee agitaattel titteeeatt igggeleetg 1500 accgccttgc catgctgctt ctaagagggg ttatctatac tcagagggcc tagggcagac 1560 1620 tttgtggccc aggetggagt gcagtggtgc gatcatagct cactgcaacc ttgagctctt 1680 gggctcaagt gattctcccc tctcagcctc ctgagtggct gggaccgcag gtgcacatca 1740 ccatgcctgg gttlttttct tlctttcttt ctttctttt ttttttggta gagatgaggi 1800 attgttacat tgcccaggct ggtctcgaat tcctgacctc gggcgatcct cctaccttgg

cctcccagtg	ctgggattac	atgtgtgagc	caccactcct	ggtctgggct	tttttttt	1860
ttttttagga	gaatgtaagg	catttgttta	gtgtttactg	agcatctact	gcatgccaag	1920
ctctgctcag	gaggccaaga	gagcaggata	ggagtgagcc	ctgcacctgg	ggtttggcct	1980
ggctggggag	ccctagcctc	aggaagcagg	actgccctgg	accccaggag	gccctgctc	2040
aacctgcgac	aagactgaaa	ggctccccag	atgcactgtg	acggtcaagg	gagagaaatg	2100
ggtgacctgg	gatcctggtg	gcttcagatc	tcattggggg	aaggagcaag	ggaccccagg	2160
ggtcagaggc	attggggggg	tagggctgtc	agagggcacc	cagcagagtg	actcacaggc	2220
tatgaggaaa	gaagtctcct	cttgcttgta	tcttcaggat	gccaagacct	gcagcccagg	2280
cctggggtgg	tggttatccc	ggagcggcct	taccaggtgc	acagctggcc	aagggaggct	2340
tggtcatggg	cagagagggc	aggaggaagg	cccttgctcc	accaccccct	ctcccaaggg	2400
agtcctatgc	ccagaactta	ccaagtagca	agactagagc	aaccagcaga	agacccttgg	2460
gggcagccat	catg					2474

<211> 2211

<212> DNA

<213> Homo sapiens

<400> 4658

60 atgttgtgag aaagtgctgg gctagctgac tcggatcatc tcctagagtt taggagaaat 120 aacaatcaca aagttigaat tigcaaaacc tilaggetti getggcagga gagaaaatac 180 cacttttgga tcataaaatg gatatggaat acatgggctc aggtaaactg ggccaaatat 240 tgaagcagat ttggagacaa aaccagatgc taaagcaggc actcttgggt gatgaccttt gtgaggggc tggtggggcc acttggatgg ccactgtggt ctttctgggt caggaactca 300 360 geagtgeeet teaccagetg etggageaea ceteaggeae eetgegttee acetgeeage agctgcatgt gctgcttgac aaagagaatc aatgtgtctc gagaaaaggt atcttgatgt 420 480 getgagtgat gitaetggee eccaagigte tigitatati acageaceat cataigtiet 540 acaacaacta gaatgeegga taataaatca catgagttet ttaatagtgg gtgataatga 600 agagttagtt agcaacgtca taactattga atgctcagat aaggaaaaga gagttccatt 660 tccaataggc attgcaattc catttactgc acgttacaga ggaaattaca gagatatcat 720 ggtgaaagtg tgtgacataa accttcaatc aagttaccta aacccaaatt cactagaagg aatgaaggga ggttataagg ggacctgigc iicagtaaaa gittacaaat igggtatcii 780 840 ttctgttgtg tcttgtttaa agaaagagte gttcacagta acaaagaaag gcctcgctct taagtcaagc atggattccc gaatatcctt aaattaccct ccaggagttt ttacctctcc 900 960 agtgctggtg cagttaaaga tccaaccagt tgacccagct ctggtggcac atttaaaagc

acagcaagat	actttctact	cagtccaatc	cacaagccct	ctgattcaca	ttcagcaccc	1020
atcaacttat	ccttttcaga	agccagtcac	tttgttttta	ccttgttctc	cataccttga	1080
taaaaacaac	cttggttctg	agatagatca	taaaagaaga	gcaagtgcca	caataaatag	1140
gattacacct	tcgtatttca	accggacaaa	aattgcctcc	ataagaaaac	ctaggaaaaa	1200
tgccagtgaa	tgtttgaaat	tactgggatt	cagaagccaa	gacagtggtt	ggtgtgggct .	1260
tgatgatgtt	gtgaaaacca	tacagagcgg	cttggtatca	gttgaattgt	atgaacattt	1320
ggagaggttt	attgtacttc	acctctcttc	caccatggac	aatagtcatt	tggttacttt	1380
tgtgaaatct	ttagaggaag	ccatgctcag	caccactgcc	tgcatagtac	tgtctcacca	1440
gaaggacaat	ccacatagaa	tagctgtttt	agtggtgcct	tccaaagatt	taagccaggt	1500
gcttaaggac	ctgcacttgg	aagggtttgg	aggacctcca	gagccatctc	gtcatttcca	1560
agttcgagaa	ggagaacaac	ttcttttaag	atttactgga	aacatatttg	cttcaagcaa	1620
cgggaaggat	tatggaaaag	actacacact	tatttttcac	ttgcaaagaa	aacctaggct	1680
ggaactccaa	atcaaagaag	tggacgaatt	tggaaactat	agttgccctc	attacaaagg	1740
caccattgtc	gtttataaag	tacctaaagg	aaagatagtc	cccaacttga	atcaatctct	1800
cgtaattaat	gaaaaccatt	ctcagttgcc	aatttgcaaa	ttaccattga	aattgccaaa	1860
gcataagaaa	ttaatcaacc	atccacagag	taccaaaaga	gtttctaagg	atcctgtaga	1920
agccctttgg	gataacttgc	tccattggct	ggctgaggag	ctctcagaag	aaaatgctga	1980
gtctctttcc	tcaactctcc	ctctgcgccg	tagcaccatt	cagctcatca	aactcaagaa	2040
ccctgatgat	ctcacagaac	agatccacga	gtttctttgc	ttctggaaaa	aatcgcttcc	2100
aactttcacc	gacaaacttc	gcctcctggc	tcgacatctc	cgcaagattg	gcaggagtga	2160
tcttgcagaa	gaactcaaat	tcaagtggga	aaataaagtg	ttcactgaac	c	2211

<211> 2513

<212> DNA

<213> Homo sapiens

gaactgtggc	agcatctggc	cttctctgca	tgaccagcgt	ggtgtggcat	ggatcacaca	60
gctgggctgg	cttgccctcc	tgtgcttctg	tgcccagttg	ataacagcag	tctttgggga	120
aggaaatcac	agttatctta	cccgccaagg	ccacgctccc	ttcccatttg	gggctcacca	180
gagcctgatc	atcctgccct	gctggcctcc	tgcctatccc	attcatttca	teccaetect	240
gccccagtt	taggctcacg	gcatcacctg	ccttgcactc	tgatggcacc	gtgctccacc	300
tcccacctcc	ttcccatctt	cctacatcag	agcctgattg	agccactgcc	ctgcacctgc	360
ctttaaggcc	cgtgtttcct	acagaatagc	attcagtctc	tgacgtgata	ttcaaagccc	420

480	ccacacttgt	agggcagggg	aagctccatg	ataagaatgg	gggtgcattc	tccatagcct
540	aacaaatatg	gaaggtgttc	ggctgtcata	ctagcatggt	atacccagac	ctggtcattc
600	ctaattttct	tcattgacgt	ttttctaacc	gcacatctat	aacaaatggt	tttttaaaat
660	caggcactgt	ctgagtattt	aactcctttt	attccaaaca	tattctagtg	tcatccatct
720	ccagccttca	aaagcctttt	ccttatcctg	catgctggtc	tgcctttact	agagcctcca
780	ccccaagcag	tacccacttt	ctggctcaca	gcattggggg	acatcctact	ctgtgtacag
840	tatttttcct	ggtggtattt	gatctagtta	ctcacttaca	tttttcccct	aataactgtt
900	aggtaaggac	taaactctaa	catgaggttg	tctttctgta	catttgttag	ctttcttggc
960	aaggacgaga	tttttctgta	ggcagcaaac	tttagacagg	cctctagtca	attatcttca
1020	catgaaagca	gctgctgtgt	agtctctcct	gggggcttgc	ttaaggcatt	tgataactat
1080	gcacagtctc	cagcaccagt	tgagcaagtg	agtacgtgat	atatgtaaat	gccgtggggg
1140	gaaaagagtg	tggcctgaat	gtgaatcgac	ctgagtggag	ctgtgggtac	atcatcgtct
1200	gtaggcatta	ctggcctcaa	gaactgggga	caaggacagg	tgtggttcct	aaagttccca
1260	cttccttata	atatttgaac	gaggtgtaaa	ttaatgtgga	cccacagagt	gaataaactg
1320	gggtaacaga	gatgctctgt	tatggaagtt	gcagctctgt	agccaagctt	aacacatgct
1380	aaaagaagct	cagtggcctt	gctattactg	accaaggaaa	atccttgtta	ctcactcttc
1440	cctgtatcag	ttgttcgcag	cgcgggtatc	gaagcactgc	taatcattca	tgggcagcca
1500	ggcaaggagg	gaggattcct	gcctacagcc	cacaatgcag	tggccaccat	ttgattcgca
1560	cgcacgggcg	tacagaaata	gctgtgatcc	ggaacataag	agatgctgga	aggtatcgaa
1620	tcagcttact	tgcttaatat	cgacgattcg	ccagagtatc	gacgcagatt	tggctggcca
1680	ccatgggctg	acaaagaaaa	gaagatcaga	gaaaaagctg	agcgtttgca	tacagggtcc
1740	gattcagaag	atgtggaaaa	cgagctgggg	ggctgctctt	tgactagcct	gtggagaagc
1800	gaaggggaag	attacgagga	cacaggcgaa	agcagccact	aactagaaaa	ctggaagcag
1860	tggaaatatg	tgagacggga	tgcgggcgtg	agaggtcagc	atgctgtgga	agatacaggg
1920	atttctgcag	tatgaagtgg	cagcttttcc	tgagcgctgc	gtgaaaaaga	cggactctct
1980	ttaggaaagg	agcctacctt	taccaaaggg	caacttccat	attitgctat	agcaaatcag
2040	gaaatgcaac	aataatgaga	ggaggaggtt	ctaggggaag	ggattagagt	cgagatcaga
2100	agccctggct	aggcccaaag	aaatgatgac	aagttttgca	tctttctttc	ataggttttc
2160	taaggtctat	cacttgtcct	ggtctcctat	ggatgacctt	ctttcttcca	ggctgcctcc
2220	ttcatctctg	gggtagggtt	tgcaaagcta	tgaacagcga	tgtcttcccc	gccctggaaa
2280	tcttgccttt	tagattcaaa	ccagatgccc	agcataaaaa	aagagcagtc	gtgtagtgaa
2340	ctcatataaa	ttttagtttc	cctctctgaa	agtcccttac	ccttggctaa	ggctgggcag
2400	catcagatga	agaataaaat	aattgttgtg	ggactgacag	taacaaaaaa	aaaatgagaa
2460	aaaatgagaa	cctcctatat	attttagttt	accataatga	ccaaatgcct	aatacgttag
2513	aat	ggtcagatga	agaataaaat	aattgttgtg	agattgacag	taacaaaaag

```
<210> 4660
<211> 4033
<212> DNA
<213> Homo sapiens
```

(100) 1000						
ggatagtcgc	cgtcaggcac	tgaggaggcc	gagagatgag	gccggagccc	accaagttct	60
gggaagttct	ttctgacaca	agtaagataa	catctctgcc	atgcccacag	gtgctgagaa	120
caagagaaat	caagtgaagg	aaggaggcgg	agtttccaag	acttgggtgt	catcatttct	180
ggggacatcc	ttgattggag	attgaagttt	ttgaaccgaa	atttagagct	gattcagaag	240
agacaaatac	agaacgtcca	agttagcagt	cgtttcccaa	atttaggaga	caatgatgca	300
ggcccaggaa	tecctaacac	tggaggatgt	ggctgtggac	ttcacctggg	aggagtggca	360
gttcctgagc	cctgctcaga	aggacctgta	ccgggacgtg	atgttggaga	actacagcaa	420
cctggtggca	gtggggtatc	aagccagcaa	accagatgca	ctctccaaat	tggaacgagg	480
agaagaaact	tgcacaacag	aagatgaaat	ctactctcga	atctgttctg	aaatcaggaa	540
aattgatgat	cctctgcagc	atcacttgca	aaatcaaagt	attcagaaga	gtgtgaaaca	600
gtgccatgaa	cagaatatgt	ttggaaatat	tgttaatcag	aacaaaggtc	atttcctgct	660
gaagcaagat	tgtgatacgt	ttgacttaca	tgaaaaaacct	ttaaaatcaa	atttaagttt	720
tgaaaaccag	aaaaggagct	ctggcctaaa	gaactctgct	gagtttaata	gagatgggaa	780
atccctttt	catgctaacc	ataaacaatt	ttatactgaa	atgaagtttc	ctgcaattgc	840
aaaacctatt	aataagtccc	agttcattaa	gcaacagaga	actcacaaca	tagagaatgc	900
ccatgtatgc	agtgaatgtg	ggaaagcctt	cctcaagttg	tctcagttta	ttgatcatca	960
gagagttcac	actggagaaa	aacctcatgt	atgcagtatg	tgtgggaaag	ctttctccag	1020
aaaatccaga	ctaatggacc	atcagagaac	tcatacagaa	ctgaaacatt	atgaatgcac	1080
tgaatgtgac	aaaaccttcc	tcaagaaatc	acagctcaat	atacatcaga	aaactcatat	1140
gggagggaaa	ccttacacat	gtagccaatg	tgggaaagcc	ttcatcaaga	agtgtcggct	1200
catttatcat	caacgaactc	atacaggaga	gaaaccccat	ggatgcagtg	tatgtgggaa	1260
ggccttctct	acaaagttca	gtctcactac	acatcagaaa	actcatacag	gagaaaaaacc	1320
ttatatatgt	agtgaatgtg	gaaaaggctt	cattgagaag	aggcgtctta	ctgcacatca	1380
tegaacteat	actggtgaga	aaccctttat	atgcaataaa	tgtgggaaag	gcttcacctt	1440
gaagaacagt	cttatcacac	atcagcaaac	tcatacagga	gagaaattat	atacatgtag	1500
tgaatgtgga	aaaggctttt	caatgaagca	ctgtctcatg	gtacatcaac	gaactcatac	1560
tggagagaaa	ccttataaat	gcaatgagtg	tggaaagggc	ttcgctttga	agagcccact	1620
catcagacat	cagcgaacac	atactggaga	gaaaccctat	gtatgcaccg	aatgtcgaaa	1680
aggtttcacc	atgaagagtg	acctcattgt	acatcagcga	actcatactg	cagagaagcc	1740

atatatatgc	aatgattgtg	gaaaaggctt	cactgtgaag	agccgcctta	ttgtgcatca	1800
gcgaactcat	actggagaaa	aaccctatgt	atgtggtgag	tgtggaaaag	gctttccagc	1860
aaagatccgg	ctaatgggac	atcaacgaac	tcatacagga	gagaaacctt	atatttgcaa	1920
tgagtgtgga	aaaggcttca	ctgagaagag	tcatctcaat	gtacatcggc	gcactcatac	1980
aggagagaaa	ccctatgtat	gcagtgaatg	tggcaaagac	ttaactggga	aaagcatgct	2040
cattgcacat	cagcgaactc	atactgggga	gaaaccttat	atatgcaatg	aatgtggaaa	2100
gggcttcacc	atgaagagta	ctctcagtat	acatcagcaa	actcatactg	gagagaagcc	2160
atacaaatgc	aatgaatgtg	ataaaacctt	caggaagaag	acatgcctca	tacaacatca	2220
gcgatttcac	acaggaaaga	$\tt cttcctttgc$	atgtactgaa	tgtggaaaat	tctctttgcg	2280
caaaaatgat	cttattacac	atcagagaat	tcacacagga	gagaaaccgt	acaaatgcag	2340
tgactgcggg	aaagccttca	ctacaaaatc	agggctcaat	gttcatcaaa	gaaaacatac	2400
aggagagagg	ccctatggat	gtagtgattg	tgggaaagct	tttgcgcact	tgtctatcct	2460
tgttaaacac	aggagaattc	acaggtagtc	attttgggaa	agcctcttgc	cagatgtagg	2520
cccttaagat	atctgcaaag	aagagtaatt	tcatgaatgc	agactacatg	gttgtttatt	2580
tagtgatcag	ttacttcatg	ttttgtgtca	gagaaaacat	gtacaaaaca	tttgagaaaa	2640
tattttagga	cattatgtct	aaaaattgta	tactgagaaa	aatcctatga	atgtggcaga	2700
ctataaaagc	ctttggtggg	aagataaacc	ccctcagaag	tgatcataga	tcatgaataa	2760
actactaatt	cgtggaaatg	taataattat	aggaacgtct	ttgcccaaaa	ataaaacttc	2820
aacagattga	gagagttcac	actggagaaa	tactttcctc	tggcaaaccc	ctgtggcagg	2880
gttttcagta	aaatgttttg	cctcatcgtg	tgctaggaaa	taatgcagaa	aaaacttatg	2940
aaaacattca	atgtagaaga	cttttaggaa	aatatcagag	aatttaatga	aggaaagaag	3000
ccttggcaaa	atgatggatg	agagctttct	gtcacaattc	taaccttaac	agataatatg	3060
cctcatgtgc	attigggaac	aggatagctt	gagccatatt	cctagttgcc	ttgtcactga	3120
ttttatacat	tttaaattgt	gacttcctct	aatcacgaac	taaaatttaa	taitgtatat	3180
aataccggag	tattttatgc	ctgcctcatt	ttattatatg	attagctcct	gcatttcttt	3240
ggttctaagt	tigigitati	tcagaataac	taaagtactt	cagaatgaaa	ctggatacag	3300
tatattcagt	tcataagtgt	aattgaccat	acttgaatat	gtttaagttt	ataagtaaca	3360
tcaggtgtta	tgaaattaat	ataaataaga	tgtttttgta	agtttttatt	ttgaaataag	3420
tttagattca	caaaatctta	ccaaaatatt	ccagagttct	tgcccttaac	ccagettett	3480
ccaaagataa	caacttacat	aagcatagtt	cattaccaaa	tacaggaaag	tgaaattgtt	3540
acaatacigi	taactacaga	atttatttgg	aaatcaccag	ttttaacatg	catgcatttt	3600
igittgittt	ttatatgtag	ctttaagaaa	tcttaacaca	tatattgatt	catgtcacca	3660
ccaccgtaat	cgagattcag	aatgttgcat	caatgcaaac	tccctcatgc	tccttgttta	3720
tagtcacaac	ctctcctcag	ccctaaccct	tggcaatcag	aaatcttttc	atctttagtt	3780
tigicatiag	tagcatgita	taaaaatgga	atcatacagc	atgtaccctt	ttggtattgg	3840

ctatttcaac	taaagcagat	taacctgaaa	tccatccaag	ttgttctgca	tatcattgtt	3900
ctgtttttat	tggccactaa	taatctatga	tacgaatata	ccactatttg	tttattcacc	3960
cactgaagga	cattagtttt	gttcccggtt	tttcttactg	ttacaaataa	aaatcctatg	4020
aacatttgta	cac					4033

<211> 4275

<212> DNA

<213> Homo sapiens

60	ttggctgggt	ggcatgacat	ggaacaaggg	accaggetet	aataatatgt	gattcaatgg
120	acttgagcaa	tgtggatggt	ttattttttg	attcaagcct	attctttttg	ttgctggatg
180	ctgcctcata	ttccccttta	accatggtat	gtttacggaa	ggaacatitg	cttgattcct
240	ctagtctgtt	tttctaagtt	aaggcaatac	tggtggagaa	gtgggttict	ttggaagagt
300	aaggagagct	caaaacaggg	gtcatcactg	acaataaagg	gagaactttg	cttcttcaat
360	agggccacaa	aaggaatatg	gtctgtgacc	tcaccctggt	gccccgggag	tgaaggaagt
420	ccctgctggg	caaatcaccg	ctacagagac	gcctgacctt	caagacctca	ggctgtggtc
480	accctcccac	acggggctcc	atccatgttg	ccactatcat	gccgggaaaa	gacaaacggt
540	gggtcagaat	gacctgtcga	cctacagaca	atggcaagaa	atcatcatca	ticiggaacc
600	tccgggaaca	aacctcaccg	cctgttggac	agcaggacat	gtgtgtccgc	ggagcttggt
660	tgcatcagca	aagaaggagc	tcagtggacc	taaaggcgcc	tttgcttcca	tttgctgctc
720	agacccgagc	cagcacaaac	aactcagcat	atgtggactt	actcttcagg	agtcaatcaa
780	gcatgtcgag	gctttcatgg	ccttggcatt	ggaagctctc	ggcctgaaga	tctgtctgga
840	atagcctgtg	tgctcccggc	ggtggaccct	ccaccagtgg	ctggatgagc	gaccgtggtt
900	accacctgga	ttcacaaccc	tacgatcatc	gagaaggtcg	ctcaagtacc	ggacattctg
960	tcaggtgctg	catgggaggc	cgtcctccag	accgcgtggc	gcgctgagtg	tgaagctgaa
1020	cactcacgag	ctccgcctga	tggccagggg	aggaggcata	ttctgcctga	cggtcctccc
1080	catccctgat	gcttgtgtta	gaaagacatg	cccatgatct	gttctggagg	geageettet
1140	tgacctacac	ggaagtgagc	agacagcagt	catttctcaa	attccacaag	aaagatetat
1200	tggatgggaa	ttccaggccc	gaaagggctc	aggcctgctt	gacacagaca	cattccaaag
1260	aagaggtgtt	accaccttag	gatctcagac	cgggctatgg	ctgcacctga	cctgcatcag
1320	ctgagtcaga	gccctgggga	atctcacatt	ccaacaagaa	ttgcaagatt	tttgatgctt
1380	tagcacggcc	tgtggctccc	gtctggctac	caggacatet	cacaggeeta	gctgcagaac
1440	tggcccggag	gccgcgatcc	cgcacaagtg	agctgctccg	cagggcgtcc	egcaactgtg

```
gctccgccgc acgctgcgcg ccgggaagag caccctcgcc gacctgctgc tgccagtcct
                                                                    1500
                                                                    1560
cttcgtggcc ttggccatgg gcttgttcat ggtgagaccc ctggccaccg agtaccctcc
                                                                    1620
cctcagactc acacctggac attaccagcg ggccgagacc tactttttca gcagtggggg
                                                                    1680
cgacaacttg gacctcaccc gtgtgcttct gcggaagttt agagatcaag atttgccctg
tgcagattta aacccacgcc agaagaattc ttcatgctgg cgcacagatc ccttttctca
                                                                    1740
                                                                    1800
cecagaatte caggatteat gtggetgeet gaagegteea aatagaagtg ctagtgetee
ctacctgacc aaccacctgg gccacacact gttgaatctc tcaggcttca atatggagga
                                                                    1860
                                                                    1920
gtacttgctg gcaccatctg aaaaaccaag gcttggaggt tggtcttttg gattaaaaat
                                                                    1980
ccccagtgaa gctggaggtg caaatggaaa catatcaaaa cccccaactc tggcaaaggt
                                                                    2040
gtggtataat cagaagggtt ttcattccct acctlcctac ttaaatcatc taaacaacct
                                                                   2100
tattttgtgg cagcacctac cccctactgt ggactggaga caatacggaa taacactcta
                                                                   2160
cagccaccca tatggagggg ccttgctgaa cgaggacaag atcctggaga gcatccgtca
                                                                    2220
gtgtggagtg geeetetgea tegtgetggg atteteeate etgtetgeat eeateggeag
                                                                    2280
ctctgtggtg agggacaggg tgattggagc caaaaggttg cagcacataa gtggccttgg
ctacaggatg tactggttca caaacttcct atatgacatg etctttact tggtttccgt
                                                                    2340
                                                                    2400
ctgcctgtgt gttgccgtta ttgtcgcctt ccagttaaca gcttttactt tccgcaagaa
                                                                    2460
cttggcagcc acggccctcc tgctgtcact tttcggatat gcaactcttc catggatgta
                                                                    2520
cctgatgtcc agaatctttt ccagttcgga cgtggctttc atttcctatg tctcactaaa
                                                                    2580
cttcatcttt ggcctttgta ccatgctcat aaccattatg ccccggttgc tagccatcat
                                                                    2640
ctccaaagct aagaatttac agaatatcta tgatgtcctc aagtgggtct ttactatttt
                                                                    2700
tecteaatte tgtettggte aaggaetggt agaactetge tataateaga eeaaatatga
                                                                    2760
ccigacceae aacticggea tigatteeta igigagteee titgagatga actitetggg
                                                                    2820
etggatette gtgcaactgg eetegeaggg cacagtactt eteetettga gggttetget
acactgggac cttctgcgat ggccaagggg tcattctact ctccaaggca cagtcaaatc
                                                                    2880
tictaaggat acagatgiig aaaaagagga aaagagagig titgaaggaa ggaccaatgg
                                                                    2940
                                                                    3000
agacattett gtgttataca accitagiaa acattatega egettittee agaatattat
                                                                    3060
tgctgtgcaa gatattagtt tgggcatacc aaaaggagag tgctttggac ttctaggggt
                                                                    3120
gaatggaget gggaagagea egaettteaa aatgetgaat ggtgaagttt etetaaette
                                                                    3180
aggacatget atcatcagga ctcccatggg agacgcgtg gacctgtctt ctgctggcac
                                                                    3240
ggeaggegtg cleatigget actgleecea geaggatgee etggaegage tietgaetgg
                                                                    3300
tigggaacat cictattatt aciglagett acgegggatt ccaaggeagt geateceiga
                                                                    3360
ggitgetgga gaeeteatea ggegettaea eetegaagee eaegeggaea aaeetgtgge
                                                                    3420
cacctacagt gggggaacca agcggaaact ctctacagcc ctggccctgg tggggaaacc
                                                                    3480
tgacattett ttattggatg ageceagete tgggatggat eeetgeteta ageggtaeet
                                                                    3540
gtggcaaaca ataatgaagg aggttcggga aggctgtgct gcggtgctga cctcccacag
                                                                    3600
gtitiggtat caggatgatg ciggocical aaaalgagit acagaggati cocicitiit
```

ctattgattg	gaatagtttc	agaaggaatg	gtaccagctc	ctctttgtac	ctctagtaga	3660
atttggctgt	gaattcgtct	ggtcctggac	ttttttgga	aaatatataa	agcatatttg	3720
gaaaatttat	gacatgtatc	tcttggactt	catgctaagc	aactataaca	aagaaacctc	3780
aacttacagt	ggcccagaca	agacacattt	tttcctctcg	ctaaattcat	gtcagtgggc	3840
tgtgcatcat	caggccaacc	aaggccttgg	gttctttcca	ccgtgccacc	atcaccatca	3900
tcatggttga	tgttgactca	tcagcaccgt	atctacatcc	tagctcaagg	gaagtggaga	3960
aaaacaaaat	ccagcatgga	ggagtgtgag	gctctttgca	caagactggc	cataatggtt	4020
aacggcagct	tcaaatgtct	tggttctcct	cagcacatca	aaaataggtt	tggtgatggt	4080
tatacagtca	aagtttggct	ctgtaaggaa	gcaaatcaat	attgcactgt	ttctgaccac	4140
ttgaagcttt	attttccagg	aattcagttc	aagggacagc	acctgaattt	attagaatat	4200
catgtgccaa	aaagatgggg	atgcctagct	gacttgttca	aagttataga	gaacaataaa	4260
accttcttga	atatt					4275

<211> 3411

<212> DNA

<213> Homo sapiens

```
60
ctataaatac atagtaaata titataaaag cicattitta tetetgggea gaaaaatagi
                                                                     120
cocagitatt iicticigca coccatiggg itatcagigi giactciagg gagaciagig
                                                                     180
actaaaaatg ataggiicia iittaaagiig atcggcaiig aaaagaaati ggacigaata
gictcactcc aacaaccaic aagagictii giggcattii cagaatigii galgicalcg
                                                                     240
tatgttttaa tggctttaaa aaaaaataca tttcaccttt tatagtaatg ttaattctat
                                                                    300
                                                                    360
gtatticcig icaatagaaa ictaattiai atticgciaa acccaagcia aaaatatata
ttgactgtga aatgtgcatt aaattgttat tgatcatagt tatcaacatt atgctatgtt
                                                                    420
                                                                    480
ataacagetg titgagatgg agaacgttit tactgigtea teagatecte atccetetee
                                                                     540
tgeageeeet eetteacitt etttaeeett aletteatet telaeeteat elgggaetaa
                                                                     600
aaaacaaaag aggaccccaa cttatcagag alciglaagi caggaaagca gicatcicag
                                                                    660
cttacctigc itigigitaa gagcatgacc taggatigcc actcailcig giigaaatat
                                                                     720
ctigocgact cacciletaa etteoogaat cattilitie accileacie tigaaagata
                                                                     780
acticagiag attagggict tactgeceaa agtatateta agaateagaa geattattat
                                                                    840
cacctiggat citaataaag aiggaaaatt toaaatoocc atacctacig agioigatii
                                                                    900
tatattttaa etteteattt gagaageeet ggillaeage eagacettet iggilligaat
                                                                    960
ctictactat ilgaciitgg caagicatii aacticicig igicigitti cicatiggia
```

aaatgaggcg	aatacaatac	tttcttgtgg	gcttttgtga	aagttaaatg	agttaatata	1020
tgttaaacac	agaacagtgg	ctagtgtaga	agaattagga	cgtaaccacc	tacaggtgtg	1080
gtggctttga	tttacgtgat	ttgaccatca	cacatatcgt	tgactcaaac	tttggtttat	1140
ctttctagaa	tttgttactc	ttttttccta	gtgaaaatat	atccttttta	ttaataatgc	1200
ccaacattcc	acttaaatgg	aaatacttat	aaatcagatt	ttcagtgaaa	gggtctaaaa	1260
cataattgta	atgatttaga	aatgtcatta	ctgtggtaaa	ttatagacct	aggctttgct	1320
aaggatttcg	tcctgttttg	tgcttcactt	atagtttttg	ttgttttact	caaagaaggg	1380
aaatggaaca	tccaatgtga	tagtattta	tgacctaggc	agattccatt	atattgcttt	1440
taaatagctc	ctatttcact	aaagaatggg	aacgaagtga	tagttacaga	tttcatcagt	1500
atgtaaaaag	tgaagggttt	cttttaggta	aagatttctt	taaatgtacc	tgtgaaatga	1560
ttcagcattt	ttaaaatgga	aatgcttttg	ctgtcttgga	gttttgttgt	ttcagaagtt	1620
ttctactgca	ctattttgga	tagtctttca	ttaaaggacc	taagcatgaa	ttactctgaa	1680
attttctagt	ttaagtattg	catagaaagt	tcattttatt	gtgttaatct	ctactaactt	1740
gaaatatgcc	ttccaaatgc	atgacttact	aaacagtatt	aaggtattgg	ctgaagtttc	1800
aaaatagcgt	taccaaatct	cttaagagtc	tttggtagtt	tgtggtcgct	gtctcttaat	1860
attaattacc	tttgttctga	aattgttgtt	ttagtaattt	tcacttcatg	taaaggcagc	1920
atttgctaat	gtagttgcta	taccttaata	ttaaacccca	gaaattctat	ttttgtctga	1980
taagtggaaa	ttctaccata	atttggggca	attccccaat	ataataagtt	gtttcctctg	2040
aaatgtttta	ttagctaaat	taatgctgca	gtaataaagt	ctataatctt	cccttcattt	2100
tactttttt	tgcgttttta	tatggaagat	ttagtgactt	attaaccaga	cttaatatgc	2160
tatggagata	atgtacagat	atgtatatat	tctttttctt	tttaaaacag	atgagctttg	2220
atccaaacct	tctccacaac	aatggacata	atgggtaccc	taatggtact	tcagctgcac	2280
tgcgtgaaac	tggggttatt	gaaaaactgt	taacctctta	cggatttatt	cagtgttcag	2340
aacgtcaagc	tagacttttc	ttccactgtt	cacagtataa	iggcaaccig	caagacttaa	2400
aagtaggagg	taatctgtca	gttctccttt	gtaaaaatgt	aatcacaaaa	tttgtcttgc	2460
atatcaattt	gttcatgagt	gtttttcaaa	aagttttatg	taaattaaaa	cattttggac	2520
ttttagttgt	actgtttagt	gaagaatatc	tgtataaacc	agcagaatcc	ttgaaagatt	2580
atatttgcat	atatttacaa	taaatgcttg	gaagatttca	ttgtttttcc	ccataatgat	2640
gactgtttac	ccgtaagatt	ttaagccagt	acgaatattg	aacttttgta	atgttactta	2700
tgattattct	gtcaggtact	attttgggta	gtacctgaaa	aaagtaatcc	ttagagtgtc	2760
acctcttaga	agctgggcgc	ggtggcttac	gcctgtaatc	ccagcacttt	gggaggccga	2820
ggcgggcaga	tcacgaggtc	aggagataga	gaccatcctg	getaacacgg	tgaaaccccg	2880
tetetactaa	aaatacaaaa	aattagctgg	gcgtggtggc	aggcacctgt	agtcccagct	2940
actcgggagg	ctgaagcagg	agaatggtgt	gaacctggga	ggcggagctt	gcagtaagcc	3000
aagatgccac	cactgcactc	cagcctgggc	gacagagcga	gacccegtct	сааааааааа	3060
aaaaagtcac	ctcttagatt	atatggaaat	tcattgtggc	tgacaccata	caaattattt	3120

cacaggcctg	gcatggtggc	tcatgcctgt	attcccagca	ttttgggagg	ccagggcaag	3180
tggattgctt	gagctcagga	attcaagacc	agcttgcagc	ctgggcaaca	tggcaaaaacc	3240
ccatctctac	aaaaaatgca	aaaattagct	gaggaacgtg	acatgtgcct	gtagtcccag	3300
ctactaagta	gactgaggtg	gaaggatcgc	ttgagcccgc	aaggtagagg	ttgcagtgag	3360
ctgagatcgt	gccgctgcac	tccagcttgg	gtgacaggga	gactctgtct	С	3411

<210> 4663 <211> 3371

<212> DNA

<213> Homo sapiens

<400> 4663

60 tactccctgg ctgccgtgaa aagacaaggc actgggcagt gatgagggtc cttctccatc 120 teccageeet eetggettee etcatettge tteaggetge ageatetace acaagaggtg 180 agtgtctccc tttacgggtt ttctggggct gctgtcacaa agcacaccaa ctggatggcc 240 taagacaaca gaaatgtatt ctctcacagt acgggaggat agaagtccaa aatcaaggtg 300 ttggcagggc tgtgctccct ctgaaggctc tagggaagaa tgcttccttt cttcttccag 360 cttctggtgg ccctggtatc cttggtgttc cctggcttgt agctgcaaca ctccaatctc 420 tgcctcagtc tttgcgtggc ctttttcact gggtgtcctc tcctcttctc ataaaaccac 480 catttattgg atttagggcc gatgctaagg cagtatgacc tcattttaat tcataacatc tacaaagacc ctacttccaa ataaggtcac attctgaggt tccatgcaga tglgaattit 540 600 gggagggeca ttatttaacc cactacactc teettgttga gaaaacagat cetgteetee 660 atgctccaga ccccaataca agagggaaca gggaatccca ccattactct catgccatcc ccttctgtac gtcagggtcc tggtgatcag tgggtcctct ttgatgaact tgagatagtc 720 780 cacaagacca totcaccagg aaccagagaa atccccacag tocagtagco caagccacco tgagtggcca cactccctct tcatcccatc tctgggcatt ctagtgagat clcaaagagc 840 900 aattttgaca tettgttgaa gateeaagag tattetteee attetgtget titteeeett 960 aaggactett atgaaacttt eetaatettg ateteetgee taeceagalt lgagggglgg 1020 gggtaatgge caagaaagag ggagacaaaa gcagaagaca tegcaaagga giggigteig 1080 acaccetact tectgetece cactecaace ceaagegeag actaecagaa cetetgeeat 1140 cteegatact gtgagteagg ceaaggteea agteaacaag geetteetgg acteeegaac 1200 caggotgaag accgocatga gototgagao toccaccago ogacagotot cagaatacot 1260 caagcatgec aaaggeegga egegeaeage cateegeaat ggaeaggtgt gggaggagte 1320 tttaaagaga ctgaggcaga aggcateett gaccaatgte acagateeca geetggactt 1380 gacticactg tetetggagg igggetgtgg igeteetget eeegiggiga gaigegaeee

gtgcagccct	taccgcacca	ttacgggaga	ctgcaataac	aggtggcggg	gcttggggtg	1440
		cactccgccc				1500
		gccttgccca				1560
		agccctctcc				1620
		gagccccgcg				1680
ccaaatttga	gaggccccct	tctctgcgtc	ccctccgcct	cgagcagagg	ctccttcccc	1740
catggggtcc	tgggctttgg	agccccggag	ccggcctgcc	gaageteage	gaggatcgtg	1800
ctggtttcag	gaggaagcct	gcgctgggcg	cegecaacag	ggctctggcg	cgctggctgc	1860
ccgcggagta	cgaggacggg	ctctccctgc	ccttcggctg	gacgccgggg	aagacgcgca	1920
acggcttccc	tctcccgctg	gcccgggagg	tatctaacaa	gattgttggc	tatctgaatg	1980
aggagggtgt	tctggaccaa	aacaggtccc	tgctcttcat	gcagtggggt	cagattgtgg	2040
atcatgacct	ggactttgcc	cctgacaccg	agciggggag	tagcgagtac	tccaaagccc	2100
agtgtgatga	gtactgtatc	cagggagaca	actgcttccc	catcatggta	cggccctgca	2160
gctgggcatc	tctgactagc	cccttgccca	ccctgatgta	gcagacattc	ccagcccatc	2220
agctaggatc	tctggataga	tctgggatac	tgatttggta	gtttcccatg	cctcacgtgt	2280
ccattcgttc	actcatccgt	ttattcagta	aatagacatt	gaccacccca	taccaggcag	2340
tgcactaagc	gctgaggatt	ccaaagtgaa	caggacactg	ctcctgctct	caagaacaaa	2400
gggccagtgt	gggaattgga	caagtaaact	actgatagca	gaatgggaac	actggagaag	2460
gttgtctaca	ttcccgaccc	agcagcattg	gtaaaacctg	cttattagaa	atgtaaaatc	2520
tcagttgggc	gaggcggctc	atgcctataa	tcccagcact	tggggaggct	gaggtgggca	2580
gatcacctga	ggtcaagagt	ttgagaccag	cctggccaac	atggtgaaac	cccgtctcta	2640
ctaaaaatac	aaaaattagc	ctggcgaggt	ggcacatgca	tgtggtccca	gctactcagg	2700
aggttgaggc	aggagaaccg	cttgaacctg	ggaggcggag	gttgcagtga	gccgagatca	2760
cgccactgca	ctccagcctg	ggcaacagag	caagattcca	actcaaaaaa	agaaaaaaaga	2820
aaaagaaaaa	aaaatgcaaa	atctcaagtc	tcacccaaga	tgttaggaaa	tagattctac	2880
atttttacca	agatcaacag	agattcattt	tatacatgat	aaagtttata	ctgctgtaca	2940
tcatcgtgga	ggatgacttc	aactccactc	agaaggactt	gaaacctgga	tgaatccgag	3000
tgagcagcag	ggatttgtag	ggatttccat	ccggaaggga	gaggagcttg	cagcaatgca	3060
tcttttctta	ttattcatgg	gtattacagt	tggagagtca	tttcttcact	caaagaataa	3120
atogaggota	tgtgctgtgg	tataaagagc	ataaactgtg	gaatgaagcc	aatgatctga	3180
atcctacatc	tttcccttcc	agtgtataag	ttcttaggca	acttacttaa	cttttctgag	3240
ccccagtttt	ctcatctata	aaatggaact	gaattgtcca	ccttacaggg	tttttttgag	3300
gactgagatg	agttaaagta	tgcataattc	ctggcacatg	gtaagtccaa	aataaataca	3360
tggcagctgt	t					3371

```
<210> 4664
```

<211> 3707

<212> DNA

<213> Homo sapiens

<400> 4664

60 aaaagatggc cggagcggcg gcggcggtgg ccgcgggagc agcagctgga gccgccgcgg cagccgtgtc ggtggcggct cccggccggg cctcggcgcc tccgccgccc ccgcccgtgt 120 180 actgtgtgtg ccggcagccg tacgacgtga accgcttcat.gatcgagtgc gatatctgca aggactggtt ccacggcagc tgtgttggag tagaagaaca tcatgctgtt gacattgacc 240 300 tgtatcactg tcccaactgt gcagttttac atggttcctc cttgatgaaa aaaaggagga 360 actggcacag acatgactac acagaaattg atgatggttc caaaccagtg caagctggaa 420 ctagaactit cattaaggaa ttacgctctc gagtcilccc aagtgccgat gaaalaatta 480 taaagatgca tggcagccag ctgacacaaa gatatctgga gaaacatgga tttgatgtcc 540 ctattatggt cccaaaatta gatgatctag gactcaggct ccctteacct acattttetg 600 tgatggatgt ggaacgttat gtaggtggtg acaaagtgat agatgtcatt gatgtggcga ggcaggcaga cagcaaaatg acacttcaca attatgttaa atacttcatg aatcctaaca 660 720 gaccaaaagt gttaaatgtg atcagccttg aattttcaga tacaaagatg tctgaattgg 780 tggaggtccc tgatatagcc aaaaaacttt cctgggtgga aaattattgg ccagatgatt 840 cagtetttee caageeattt gtteagaaat attgettaat gggagtteaa gaeagetata 900 cagattteca cattgactic ggtggaactt cagtetggta ceatgtecte tggggtgaga 960 agatttttta tttaataaag ccaacagatg aaaatttggc acgilatgaa icitggagtt 1020 catctgtgac ccagagtgag gtgttctttg gagataaggt ggataaatgc tacaaatgtg tggtaaagca gggacatacc ttatttgttc ctacagggtg gatccatgct gtgclcactt 1080 1140 ctcaggactg tatggcttlt ggggggaact tcctgcacaa ccttaacatt ggcatgcagc 1200 tcaggtgtta tgagatggag aaaaggctaa aaacaccaga tcttttcaaa tlccctttct 1260 ttgaagccat atgttggtit giagccaaaa acitgcigga aaccetgaaa gaacigagag 1320 aagatggttt ccagcctcaa acttacctag tacagggagt gaaagcactg catactgctt 1380 taaaattatg gatgaaaaaa gaacttgtat ctgaacatgc ctttgaaatt ccagacaatg 1440 ttagacctgg acaccttati aaagaactti ctaaagiaat tcgagcaata gaggaggaaa 1500 acggcaaacc agttaaatct cagggaattc ctattgtgtg tccagtttca cgatcctcaa 1560 atgaagcaac ttccccatac cattcccgaa gaaagatgag gaaacttcga gatcataatg 1620 teegaactee tietaaceta gacateetag ageteeacae aagggaggie eleaaaagal 1680 tagagatgtg tecatgggaa gaggacatet tgagetetaa aetgaatgga aaatteaaca 1740 aacatctcca accatcctcc acagtacctg aatggagage gaaagataat gatctacgat 1800 tactgctgac aaatggaaga ataattaaag atgaaaggca gccctttgca gatcaaagtc

tttatacagc	agatagtgaa	aatgaagagg	ataaaagaag	gacaaaaaaag	gcaaaaatga	1860
agatagaaga	gagttcagga	gtagagggag	tggaacatga	agaatctcaa	aaaccactga	1920
atgggttttt	tacacgtgtg	aaatcagaac	tcaggagtag	atcatcagga	tattctgata	1980
tttctgagtc	agaagactcc	ggacccgagt	gcactgcact	gaaaagtatc	tttaccactg	2040
aagagtctga	aagttcaggt	gatgaaaaga	aacaagaaat	aacatccaac	tttaaggagg	2100
aatctaatgt	gatgaggaac	ttccttcaaa	agagccagaa	gccatctaga	agtgaaattc	2160
caattaaaag	ggaatgtcct	acctcgacga	gcacagagga	agaagctatt	cagggcatgc	2220
tgtctatggc	agggttgcac	tattccacgt	gtttacaaag	gcaaatacaa	agcacagact	2280
gcagtggtga	aagaaactct	ctccaggatc	ccagcagctg	ccatggcagt	aaccatgagg	2340
ttaggcagtt	gtatcgctat	gataaaccag	tggaatgtgg	ataccatgtc	aagactgaag	2400
atccagactt	gaggacttcc	tcctggatta	aacagtttga	tacttccaga	tttcatcctc	2460
aggatctaag	tagaagccag	aaatgcatca	gaaaggaagg	ttcatcagaa	attagtcaga	2520
gggtacaaag	taggaattat	gtggacagca	gcggctcaag	ccttcagaat	ggaaagtata	2580
tgcagaattc	aaacctgact	tcgggggcgt	gccagataag	taatggcagt	ctaagcccag	2640
aaaggccagt	tggtgaaact	tccttctcgg	tgccccttca	ccccaccaag	agaccggcat	2700
caaatcccca	cctatcagca	accaggcaac	aaaaggtaaa	cgtccaaaaa	aaggaatggc	2760
aacagccaaa	caacgtcttg	ggaagatcct	taagttgaac	agaaatggcc	atgcacgttt	2820
ctttgtgtga	cagagctgct	gttgcagcca	ttcttccctt	tggagaccag	tctaggggtg	2880
caggageetg	gagcttccgc	tgtcccctg	cctggagcag	tttgtgtgta	tagtaagaac	2940
actgcccgaa	gaacagaatg	aacctgatgc	tgcattttca	ctgtgccaca	cccactcagc	3000
aataaccatt	ttggacctgg	tgggggagag	gaagaaggag	ggtagaacct	taaaaagaga	3060
ccttgaactg	gaaagggtct	cttgtcaggg	cttgaatttt	attttgttgt	tggtagtgtc	3120
ttgatgtatt	ttcagtggta	gggtaaagaa	ttatcaataa	tttatttaac	agatttttt	3180
ttaaagttaa	cagcitttaa	attcttttt	taaagctatt	tatttggaag	atttctggag	3240
aaatatctca	ctaatttaga	tgtaagaatg	tgaaggtttt	taaattattt	ttgatagtgt	3300
gtgtgttaca	tgtggggaag	ggccacagta	acagtaacta	gtctggactc	ttaaatttga	3360
tattcaggtt	aaagtcttaa	acagggattt	gatgcattaa	ttattttaaa	ttaagatgta	3420
tatgaaaatc	attttatttt	atatatttca	tgtgtttttt	ataagctatt	agcttcgctt	3480
ttgctaacat	ccaaggtgca	tactgttatc	caggttgatt	accttatatc	ccaccttccc	3540
tctgcactcc	ccatcatttt	gtgatgaccc	agtaagactc	ttctctttgc	agggaaacac	3600
tttcgtagcc	aatgtgtaag	aactccatga	aagateeete	atttctcatt	tcgtttgaca	3660
ttgtgatttt	cttctcaaca	ttaaaaaaaaa	taggcttttg	cattttc		3707

<211> 3660 <212> DNA

<213> Homo sapiens

60	gacccgggac	ggcgccgctg	gcgaccgtga	gcatggagcc	gcagtcccgg	ttccggggtc
120	tgcagctgga	gtcccacgcc	cccggtctgt	cgccccacgt	gtccggccgc	gacctgccca
180	gctgagcggt	cctgaggtca	ccctgccct	ttagaaggca	tctggaccct	atggaggctc
240	tggagaagct	caccggaagt	acaaggagaa	ctgcgcctgg	ggttaagaaa	taatgcggaa
300	cacagcgaga	ctacgacttc	tggccacctc	gctgagagga	ttccgagggg	tctcgctgaa
360	ggcggcaaga	ctgccgcgac	gcctcttcca	gccagcaaca	cctcttccag	gtggcctctt
420	tcagggcccc	gacccagtgc	tggaaatcaa	atgaaaccgc	ggtgtcccct	acggcttcat
480	aataacagcg	ctccttcatc	ctgccttctt	cctgccgacc	caaaatctgc	ggatggaccc
540	tgccaccaag	gctgaccttc	aggagcggcg	gagacaggcg	ggccaacatc	acctgtgggt
600	gtcatacagg	ggccaccttc	ctgcgggtgt	gaccccaagt	tgtcctggat	gtttatccaa
660	gaaggttcag	agcctcctgg	ggtgccccac	gggtactggt	ccgcttcact	aagagttcga
720	gtggaggtca	tgagtccgag	aggaagtcga	atcctgtatg	gacgctgcga	agggcctcaa
780	taccccagga	ctcgtatcgg	ggaagacgga	ctagaagaaa	ctctcctgcg	ttcacgtccc
840	gacagccagg	gttccagact	aactggctga	attgccttga	gaatcccaag	caggcagcaa
900	ctgttcccga	cttcagctcg	tggtgcagcc	gagaaggagc	ctcgacccag	gcaagatcgt
960	tgggccatgt	caaatacgcc	cccgggatgg	gccgggtgga	catcgccagg	aggtggagta
1020	ctgttcatcc	cccccggcc	tcgtcctcct	tggctccagc	gccccagcag	tectggaccg
1080	aggaatgtcc	agctgtcccc	cctctgccag	cagcggctag	gaatgaggag	cgagcacaga
1140	gacatcttct	caatgttcat	acgtctggat	gaggtcacca	ggtgtacgag	agccgtatgt
1200	aatgaatgca	tctccgcgcc	agctctgctt	ggagaggacg	ccaatcagag	atcccttccc
1260	ggctacgatt	aaaatcccag	ccgccgtttt	tacaaagtca	ctgccatttg	agaccggctt
1320	gaagagattg	ccccattaag	aatttaagtg	ggggaagatg	cttcagcccc	ggagtgagcc
1380	tgggtcaatg	ctccaagatc	cgaggcacgg	gaggttttgg	cggtgaatgg	ctctgaccag
1440	caccacctct	gccgctggag	ccaaggacac	ttccagggca	gctggtgtac	aggagaccaa
1500	ggcttctccc	caccacgccc	tcgtacgcct	gccggcgaga	ctatgaggcg	acgtggtcag
1560	agcgtgagca	ccactacagc	tgttcgtcag	aacttcgaca	catgagccag	atagctgctc
1620	ctgcacaagc	cgacgacccc	gcggccccga	tacaagctga	cgtgcacgtc	cgccgccctg
1680	tatgttcctc	cccccggat	cagccagctg	atgatggagg	ctgggctagc	agccccgctt
1740	atctacaagc	ctacggcatg	atgtgcggct	acgcgctcgg	ccatttccac	cagagatett
1800	ggaggccccc	ctttgtatat	ccaccgtcct	aagaagcacc	gcagccaggg	cccacgcctt
1860	aacacactgg	cttgcggctc	gcatcaagta	tccttcaaag	ggtgaataac	aggtgcagct

cctccctggg	ctacgccgtg	gttgtgattg	acggcagggg	ctcctgtcag	cgagggcttc	1920
ggttcgaagg	ggccctgaaa	aaccaaatgg	gccaggtgga	gatcgaggac	caggtggagg	1980
gcccgcagtt	cgtggccgag	aagtatggct	tcatcgacct	gagccgagtt	gccatccatg	2040
gctggtccta	cgggggcttc	ctctcgctca	tggggctaat	ccacaagccc	caggtgttca	2100
aggtggccat	cgcgggtgcc	ccggtcaccg	tctggatggc	ctacgacaca	gggtacactg	2160
agcgctacat	ggacgtccct	gagaacaacc	agcacggcta	tgaggcgggt	tccgtggccc	2220
tgcacgtgga	gaagctgccc	aatgagccca	accgcttgct	tatcctccac	ggcttcctgg	2280
acgaaaacgt	gcacttttc	cacacaaact	tcctcgtctc	ccaactgatc	cgagcaggga	2340
aaccttacca	gctccagatc	taccccaacg	agagacacag	tattcgctgc	cccgagtcgg	2400
gcgagcacta	tgaagtcacg	ttgctgcact	ttctacagga	atacctctga	gcctgcccac	2460
cgggagccgc	cacatcacag	cacaagtggc	tgcagcctcc	gcggggaacc	aggcgggagg	2520
gactgagtgg	cccgcgggcc	ccagtgaggc	actttgtccc	gcccagcgct	ggccagcccc	2580
gaggagccgc	tgccttcacc	gccccgacgc	cttttatcct	tttttaaacg	ctcttgggtt	2640
ttatgtccgc	tgcttcttgg	tigccgagac	agagagatgg	tggtctcggg	ccagcccctc	2700
ctctccccgc	cttctgggag	gaggaggtca	cacgctgatg	ggcactggag	aggccagaag	2760
agactcagag	gagcgggctg	ccttccgcct	ggggctccct	gtgacctctc	agtcccctgg	2820
cccggccagc	caccgtcccc	agcacccaag	catgcaattg	cctgtccccc	ccggccagcc	2880
tccccaactt	gatgtttgtg	ttttgtttgg	ggggatattt	ttcataatta	tttaaaagac	2940
aggccgggcg	cggtggctca	cgtctgtaat	cccagcactt	tgggaggctg	aggcgggcgg	3000
atcacctgag	gttgggagtt	caagaccagc	ctggccaaca	tggggaaacc	ccgtctctac	3060
taaaaataca	aaaaattagc	cgggtgtggt	ggcgcgtgcc	tataatccca	gctactcggg	3120
aggctgaggc	aggagaatcg	cttgaacccg	ggaggtggag	gttgcggtga	gccaagatcg	3180
caccattgca	ctccagcctg	ggcaacaaga	gcgaaactct	gtctcaaaat	aaataaaaaa	3240
taaaagacag	aaagcaaggg	gtgcctaaat	ctagacttgg	ggtccacacc	gggcagcggg	3300
gttgcaaccc	agcacctggt	aggctccatt	tcttcccaag	cccgagcaga	gggtcatgcg	3360
ggccccacag	gagaagcggc	cagggcccgc	ggggggcacc	acctgtggac	agccctcctg	3420
tccccaagct	ttcaggcagg	cactgaaacg	caccgaactt	ccacgctctg	ctggtcagtg	3480
gcggctgtcc	cctcccagc	ccagccgccc	agccacatgt	gtctgcctga	cccgtacaca	3540
ccaggggttc	cggggttggg	agctgaacca	tececacete	agggttatat	ttccctctcc	3600
ccttccctcc	ccgccaagag	ctctgccagg	ggcgggcaaa	aaaaaaagta	aaaagaaaag	3660

<211> 2235

<212> DNA

<213> Homo sapiens

<400> 4666 60 gaggaggagg aggagcaggc gccgccatgg ccgccgctat caccgacatg gccgacctgg 120 aggagetete cegeetgage cetetgeece eeggeageee gggtteggeg gegeggggee gggctgagcc ccccgaggag gaggaggaag aggaggagga ggaagaggag gcggaggccg 180 240 aggcggtggc ggcgclgctg ctgaacggcg gcagcggtgg gggcggcgga ggcggcggcg 300 gaggagtggg gggcggcgag gcagagacga tgtcggagcc gagccccgag agcgccagcc 360 aggccgggga ggacgaagac gaggaggagg atgacgagga ggaggaagat gagagcagca 420 gcagcggcgg gggtgaggag gagagtagcg ccgagagcct ggtgggcagc agcggcggga geageagega egagaeeege tegttgagee eeggegeege cageageage ageggggatg 480 540 gggacggcaa ggagggcctg gaggagccca agggaccgcg gggcagccag ggcggcggcg 600 ggggcggcag cagtagcagc agcgtagtct ccagcggcgg cgacgagggc tacgggactg 660 ggggaggcgg aagcagcgcg acctecgggg gecggegggg cagettggag atgtegtegg 720 atggggaacc cctgagccgc atggactcgg aggacagcat aagcagtact ataatggatg 780 tagacagcac aatttccagt gggcgttcaa ctccagcaat gatgaatgga caaggaagca 840 ctactictic aagcaaaaat attgcctata attgttgttg ggaccagtgc taggcttgct 900 tcaactctag cccagatctg gcagatcaca tccgttccat acatgtagat ggtcagcgag 960 gaggggtatt igitigetta iggaaaggit giaaagtata taacacteca telaccagte 1020 aaagttggtt acaaaggcat atgctgacac acagtggaga caaacctttc aagtgtgttg 1080 ttggtggctg caatgccagc tttgcttctc agggagggct agctcgtcat gtacccacac 1140 acticagica geagaactee teaaaagiti etageeagee aaaggeeaaa gaagaatete 1200 cttctaaagc tggaatgaac aaaaggagga aattaaagaa caaaagacga cgctcattac 1260 caeggecaca tgattiette gatgeacaaa eaetggatge gataagacat egagecatat getttaacet eteageteat atagaaagtt tagggaaggg acacagtgtt gttttteata 1320 gtactgtaat agctaagaga aaagaagatt ctgggaagat caaacttttg cttcattgga 1380 1440 tgcctgaaga cattctgcct gatgtgtggg tgaatgaaag tgaacgacat cagttaaaaa ctaaagtagt tcatttatca aagctaccca aagatactgc cttgcttttg gacccaaaca 1500 1560 tatacagaac aatgccgcag aagaggttga agagaactcl gataagaaaa gtgttcaatt 1620 tgtatttaag caaacagtga acgacgtttg caatcaacta aaaattcgtc tatcgaatta 1680 gggctgaaaa ttactgttaa agagtgttgc agtatgtctg gtggctccct tttcaggact

agggetttet eatggagtae agtatgitaa tatitaeeta tataaetaat etgitaaegg

tttttgaaaa acctitcaaa ttatttgaat aatcticata ttttcattta acctatatga ctctaatttt ttttctgagg aaatcatttg giititgagt tgitttttct taatgtaaga

aaaattgtat ittittaca agiateitea aactgaatei titatgeace aaagilggte

ttgaaaagga aaataaaatc actttettge ttggtaagca agaagccata tegattttt

ttaacttaca gaaatggaaa tatgtgtaac tigitagtat tgtattaaac aaatgttgca

17401800

1860 1920

1980

2040

tagagataat	agaacattgc	ttgtaaataa	ttcagcagat	ttgtaatata	ttttatatt	2100
ttgaaatgta	ctgtagatgt	tttctagagg	catgaaagtt	aaatgtatat	attatggtag	2160
aaataatatt	gaaggatatt	gtacttcact	agtgctgcca	gaggaattgt	taataaaagc	2220
accttcttta	acaat					2235

<211> 2526

<212> DNA

<213> Homo sapiens

<400> 4667

tigitaatgc tgaaaaacgt gggtttttca ttgaggaaag atgatgcttt atcacttaag 60 120 gaatigatca cigaageeea gaccaaagee ageggggeag geagegagii teaggaceag accaggatic ggtttatgct agagacgatg tiggccctga agaacaatga catgcgcaaa 180 240 attccagget atgaccccga gcccgtggag aagctgagga aactgcagag agctttggtg agtcaaggaa ctttcaattc tgttttgctc agagcttccg tgtcggcatg agcctgtcat 300 360 gaaaatcaag aaatagtcag acctgcacgt cagggtggtg tctggtgagt gagagtgctc 420 agaggcagag tgaacgcggc tcggttttcc tcaggggctg tcccagtggt ctgggcgtgc 480 tegtgeagaa tgteegtgtg tggggaagee eatgttggge gagegteete tttgeeteae 540 cccgggtgct aaacttgggc tgcaccattg gatcatccga tgctttcaga gaaggccaag 600 tecceateeg gaccaaggaa attgggatga egtaggteet ttgactetet etgacatgea 660 gccaggatgg agactggggc agcctccctg tcggagtgag tctttcagcc cctgtcatct 720 caagtccaaa tcattgctga cccaggctct tccaggcatg tgctcacctg ccactgagcc tgaggcagct ctgttactct ctcctgttgg gacagctttc caaacacaaa acactgagtt 780 840 taccgctagt tttggcattt ttccaaacag gctaaccgtc ccaattgcca gtgacccatt ccagtattca tgcaccagaa atgtcatggt caaaaacaaa aagccaccgg tgatggaagg 900 960 acggaatatg aaccaattaa aagcatttca ttagctcttc tctcttcatc aggtccgcaa 1020 cgccggctca ggttctgaga cgcagcttcg cgtctcctgg gacagtgtct tgagtgcgga gcagacgggt cgctggtgga tigtggggtc cgcctggagt ggggccccga tgatcgacaa 1080 cagteaceat acgeacetge agaageaget tgtggggaeg gtagggaeae ceatgeteaa 1140 1200 ggctgccagg cagaggcacc cccctgtgtg gtgtgtgtc ctggctttac ctggagcagc tetettaetg ttettgaatg gteaetgaaa tgtaeaaggt ttatetggag geettaeaga 1260 1320 aattgctatt aatattacat tgtgatataa ttattccatt tctgttgtat ctttttattg gactttaaaa gatctaaaag ttgaatggca cggggcaggg gagcgcatga ggacctgaac 1380 1440 gctcacgggg caggagtccg ttgctgagac ccctcaggct gcccccggg ctgtgcattc

taggctacag	gctatggatt	tcacattgtg	gaaattcaag	cccatttttc	atccagactt	1500
ccctcgttat	cccctcatgt	ccttccctgt	cccagattct	gtgtcactca	tggtcacggt	1560
ttcccaggct	ctgccagtga	ctggttctct	gttttattca	tcgtcgcttg	ccctgaggtt	1620
gaaccacgac	tgagaagcct	cccacatgag	tgttttctcc	cccagcggct	gtggcttcac	1680
tatgtgttcc	ccttgcacag	aggggcgttg	cccgagagtg	gacttggtgc	catccgccca	1740
cgtagacctc	acagttctat	tcagtgacac	tgaagtacag	tttggcacct	ctcgggaggc	1800
agtttgcatg	gagacatgcg	tatgacttgt	ccattcatcg	tgcttccagg	tcagttcaaa	1860
gatcctagaa	ctcgcccgga	agcagaggat	gaacacagac	atccggagaa	acatattctg	1920
cacaataatg	acaagtgaag	atttttgga	tgcttttgaa	aagcttctga	agtaagcatt	1980
tgtgtgcaca	ttttgaccta	ttaatgagat	gctgtaaaat	aagtaagttg	atttgctttt	2040
tgagttttgg	acgactgagt	ctgatgctgc	ctagggagga	ctgggctgtt	ctcatgacca	2100
tccgctcacg	ctgttggatt	cgtgcatctc	tgggttagtc	gtcatgttcc	tattaatttg	2160
cctctagtag	ttcctcatta	gaacatttta	tgatatataa	agattggggg	tttggggttt	2220
tttttttga	gacagggtct	tgccatgttg	ctcaggctgg	agactgggga	gttttgactg	2280
aaagtatctg	atgaccttgt	cagtggctgt	tggttgtcac	gaattgctta	aaaatacata	2340
aatcttcata	ataaagctat	tctgattctt	catttaaaac	atcctggtat	agtttctata	2400
attagttgta	ataagatttt	aaagatcctg	taatcccagc	tacttagaag	gctgagacag	2460
gaggatcact	ttagcccagt	agttcaagac	cagcctgggc	aacatactga	gacccccatc	2520
tcattt						2526

<211> 2510

<212> DNA

<213> Homo sapiens

acttccctcc	tccccgcgct	gcgagcagca	tcctctgcag	accctcggtc	ctcgcgcccg	60
gggtcgtccc	gctcctgcgg	ctcagcgtgg	tggcctcccc	tegecegeea	ccccggcaac	120
tttctctgcc	cgctcccgcg	ggttgggggc	tgccgtgccg	ggggctaact	gggggcagcc	180
tctggagaag	ggcttcagag	tcccggagac	gcccgccacc	cgcagcctgc	ccgcggtggg	240
cctgccgtcg	gatctcggca	ccctctcctc	ccctctcccc	gaaccatgac	cgagatgagc	300
gagaaggaga	acgaaccgga	tgacgcggcc	acccacagcc	ccccagggac	cgtctccgcc	360
ctccaggaaa	ccaageteca	gcgattcaag	cgctccctct	ccctcaagac	catectecga	420
agtaagagct	tggagaactt	cttccttcgc	tcgggctctg	agctcaagtg	ccccaccgag	480
gtgctgctga	cgcccccaac	cccactgccc	cctccctccc	caccacccac	agcctcggac	540

aggggcctgg	ctaccccatc	ccctcccca	tgcccagtcc	cacgccccct	ggcagcgctc	600
aaaccagtga	ggctgcacag	cttccaggaa	catgtcttca	agcgagctag	cccttgtgag	660
ctgtgccacc	agctcatcgt	aggaaactcc	aaacagggct	tgcgatgtaa	gatgtgcaaa	720
gtcagcgtcc	acctctggtg	ctctgaggag	atctcccacc	agcaatgccc	aggcaagacg	780
tccacctcct	tccgccgcaa	cttcagttcc	cctctcctgg	tgcatgagcc	gccaccagtc	840
tgtgccacaa	gcaaagagtc	cccacccact	ggggacagtg	ggaaggtgga	ccctgtctac	900
gagaccctgc	gctatggcac	ctccctggca	ctgatgaacc	gctccagttt	cagcagcacc	960
tctgagtccc	cgacaaggag	cctgagtgag	cgggatgagc	tgaccgagga	tggggaaggc	1020
agcatccgca	gctctgagga	ggggcctggt	gacagtgcat	ctccagtatt	cacagcccca	1080
gcagagagtg	aagggccagg	accagaggag	aagagtcctg	gacagcagct	ccccaaagcc	1140
accctgcgga	aggatgtggg	gcccatgtac	tcctacgttg	cactctacaa	gtttctgccc	1200
caggagaaca	atgatctggc	tctgcagcct	ggagatcgga	tcatgctggt	ggatgactct	1260
aacgaggact	ggtggaaggg	caagatcggc	gaccgggttg	gcttcttccc	agctaatttt	1320
gtgcaacggg	tgaggccagg	cgagaatgit	tggcgctgct	gccaaccctt	ctccgggaac	1380
aaggaacagg	gttacatgag	cctcaaggag	aaccagatct	gcgtgggcgt	gggcagaagc	1440
aaggatgctg	acggcttcat	ccgcgtcagc	agtggcaaga	agcggggcct	ggtgccagtc	1500
gacgccctga	ctgagatctg	agaggagcca	agggaaccca	gatgacaccc	ttgcccatgc	1560
ctggaccttg	cttctggcca	gggaggggat	caggccccct	tgtccactcc	atacccttcc	1620
tccctctgtc	cctctcctag	gtgccactta	ccgtggctta	ggagcctttt	gtactgggga	1680
agattttatt	tcttgggtgg	ggtgccctga	gtgggttgat	cctctgggac	ttggcgggga	1740
tggggtgggg	tgggagggaa	tgaggaagct	acaggtaggt	ccacccaacg	cccaggcacc	1800
ccagtctact	tgctgggctg	agtccagccc	tggggaagat	tcctggggaa	tttatttgct	1860
gcttctccca	gccttcccct	gcccacacca	cctgcgcatg	cctggcacac	ccctctcccc	1920
acagageete	tgcttgggtc	tatgtgtgtg	tgaccgtgca	tctgctcctt	tgcaagtggg	1980
gtcttgggag	caggcctttt	tgtgtcctcg	gccctccacc	cttggtgggg	aggggaccca	2040
ggaccatagg	aaggtcctga	gtccacctcc	tggtctccac	ctctgtcatt	ccgtcaattc	2100
tcaggaaagt	tttgcaggaa	tctctcccgt	tacttgaaac	ctgggcggac	agatgagggg	2160
aaggctgagg	tccctggaag	gtgtaacaaa	tcagagacta	tctgttcaag	tagaaaaccc	2220
aagagctggc	caggtgtggt	ggctcatgcc	tgtaatccca	gcactttggg	aggctgaggt	2280
gggcagatca	cctgaggtca	ggagttcaag	accagectgg	ccaacatggt	gaaaccccgt	2340
ctctactaaa	agtacgaaac	ttagccaggc	atgatggggg	tctgcctgta	atgccagcta	2400
ctccagagge	t gaggcaaga	gaatcacttg	aacccgggag	gcggaggttt	cagtgagccg	2460
agattgcacc	attgcacccc	agcctgggca	acagagcgag	actccgtctc		2510

<211> 2536

<212> DNA

<213> Homo sapiens

```
aagttggagt geagtggeac aatettggtt caetgeagee tetgeeteec aggeecaage
                                                                     60
                                                                     120
catcttccca cctcagcctc ctgagtagct gggattacag gcacacacca ccacacctgg
                                                                     180
ctaatttttg tgtcgttttt tttctgtaga gacgaggttt cgccatgttc ccgaggctgg
                                                                     240
tettgaacte atgageteaa geggtetgee ggeettggee tegeaaagtg etgagattag
aggogtgage egecteacte agectteatt cetttttatg gecaaataat attetgttgg
                                                                     300
atgaatgggc cacgttgtgt ttttccattt atccgttgac aggcactttg ttaccacttt
                                                                     360
tiggcatite tiegaggate illalagiet gaeagitiga titteatiig gigaeaaggg
                                                                     420
                                                                     480
gatigitett acacaaggee tiggtegiae tgatgetgte agetggagga tgiattitge
ttcctgttca cattgtctgt gcgtgggatg gtcatgagtc agggggtaga gcgggtgtgt
                                                                     540
                                                                     600
tagccttggg glattitggt gagacaattc ggtttaagga aggaaacacg ttttcaggac
                                                                     660
ccgtacccca aaacacaacc ataacccgag gtcgccactt ggctgagctg tcccccacag
                                                                     720
ggacticagg tecetggite etecateece aatetgette ettecatiee egecagaace
                                                                     780
agggtatttt ccaagggcag gtttcacatc cacttggggc agggatctta agccacagta
                                                                     840
agactaattt ccaggtccca ggtctcctgg gaaaccccaa aagactgagc cctgccgagt
                                                                     900
gagaacatte tetgaagete tgataateaa aacagegtgg caceteacaa geeteageag
                                                                     960
cccagaggeg gaccacatgt ataaaataac gccatgtgta gtatttaaca aaggaggete
                                                                    1020
cccaatccag ggagggaagg attettteac aaaaggeatg aaaacagttg ettecatgtt
                                                                    1080
tggaagcagt titgtiitgi tittigtiig tilltitgat acatggicli gcicigiggc
ccaggetgga gigcagigge aigalcacag eleacigeag celiggeele aigggeleaa
                                                                    1140
                                                                    1200
gcaatccccc tgcctcggcc tcccgagtag ctgggattat agatgtgcac caccacgccc
agctaatttt ttaatttiit atagagaigg cgtcttgtgg ccaggcacag tggctcacac
                                                                    1260
                                                                    1320
ctgtaatccc agcactiigg gaggccgagg taggagggtc actigaggic aggagtiiga
                                                                    1380
gaccagectg gecaacatgg tgaaaceeca tetetactaa aaatacaaaa attagetggg
                                                                    1440
tatggtagtg ticaggeeca ggtatagggg gteageecag agetgtgggt gaaaggaeag
                                                                    1500
geceatggte ceaectgtgg gageaggaga ggaceaggee gatgageett geagggggea
                                                                    1560
egecagetig iggieceage lagiticige teccaecace eccateceae teccagggag
ggatgiceet tetagggeta ecceeticee igeageeeig geigeteage ageegeeitg
                                                                    1620
                                                                    1680
ageaageeet tatecactic electggeet gggggggggg cacgeatetg ecagteetgt
                                                                    1740
gacegicoco licagececa icicigagag iggggilloc igagagicig cocileigec
                                                                    1800
cegeacacea ggggeceaga getgeagaag ecceeacagg etttgtteec ageagggete
```

agagggctcc	tggtgctcca	tcagcagtcc	ctggaggaca	cggagcagtc	tcaagtgcag	1860
ccgtccatgt	ggccttcagc	ccttgtgatt	ccatgagcta	gcatcagccg	tcccctcagt	1920
gggtggcaga	ggctgcaggg	agcagaagaa	agagcccatc	cttgggttct	cctggcttct	1980
cgctcagagc	tgctctctgc	tgtgaaggag	gaagtaggca	tatcccagtg	agittigcgt	2040
cgcctacatc	caggcagctc	agccataagt	cagcccaaag	cccgccagct	gggagctgca	2100
ctgagcccct	ggtcacaaaa	gtgaacgtca	caggacgggc	caactctgcc	caggcctcgc	2160
tgtgtgcaca	tacacaggtg	tacgcatgta	tgtaggcatg	aacatgtggg	tggacatgtg	2220
catgcaagtg	ttttgtgtac	aagcatatac	gtgtggctgg	gcacatgtgt	gtgcatgagt	2280
gttcatatgg	aagtatatgc	tgggcgaggc	ggctggatca	cttgaggcta	gaagttggag	2340
accagcctgg	ccaacatggc	aaagccctgt	ctctattaaa	agtacaaaaa	ttagccagac	2400
gtggtgatgc	acgcctatat	cccagctaca	tgggaggctg	aagcacgaga	atctcttgaa	2460
cccagtaggc	agaggttgca	gtgagctgag	atcgcaccgc	tgcattccag	cctgggtgac	2520
agagcaaggc	tccgtc					2536

<211> 2486

<212> DNA

<213> Homo sapiens

<400> 4670

60 aatttageca getgtggtgg cacatacetg taateecage taettgagag aetgaggeag gagaateact tgaaceggga gatggagttt geagtgagee gagatggtge eactgtaete 120 180 cagcotgggt gacagagcaa gactotgtot ccaaaaaaaa aaaaaaaaaa aaaaaaaaga cattleaage tggaagattt ggtteectaa ettlgageet agetetttea ttaaagtaat 240 300 aataaaagta gaactctaca titatataat ggittigact itccaaagig attitcacat ctcagcagtc ctgtgaagga claaataagg tgttlcaggg tagacttggc attgtgttit 360 420 gcaaagaagg tccaaggcca tgcagctatt tggtgacaga attgaaagta aagcctgatt 480 cicilgeige aaggegacii igelaleiag aageeagggi cactagacaa gatgeagtea 540 acaaataagt etecagaaca tatgacatet ecageetaaa ecaageteae ettteeatge 600 tggctccctc atgcagacgg aggacatccg cttggagcca gatetatacg aagcctgcaa 660 gagtgacate aaaaactlet glicegetgi geaatalgge aaegeleaga ttategaatg 720 tetgaaagaa aacaagaage agetaageac eegetgeeac caaaaagtat ttaagetgea 780 ggagacagag atgatggacc cagagctaga ctacaccctc atgagggtct gcaagcagat gataaagagg tictgiccgg aagcagatic taaaaccaig tigcagigci igaagcaaaa 840 900 taaaaacagt gaattgatgg atcccaaatg caaacagatg ataaccaagc gccagatcac

```
960
ccagaacaca ggtaagatet tggettgget eteetggeee egtggagtat etgaaaagga
                                                                    1020
attcagtggc tgtagagtga cctgctcaaa ctcccagggc tttgttgcct gggaatttta
                                                                    1080
agggaggagt ctgagtgtaa gcagggcctt cctcctttga ggagcatcca gaaaaatgga
gggagagtca ggggagagag gaggccacaa gaaccagaaa actgccctaa aagaacgttc
                                                                    1140
                                                                    1200
agaaggaatc aggccggcag tccttggaaa gaaaaatcta gaaattcaat aaaacttcat
                                                                    1260
gagtgtgcca ggagaatgta cgggtaatct gattcggaac agaaacattt cacctctgag
                                                                    1320
ttggaagacc tcgtaagtta atggtcacag tgagttggat attgtatttc tttttcagtg
                                                                    1380
ttctcaaaag tgtctgttat ggggaaggtt gctgatgtcc ccttgatttt tctgaggact
                                                                    1440
ccttagagta ttggagtctg cacaaaaccc cgcagagtag aaagattcct gaggacctcc
                                                                    1500
agaagtactc gttaacaagt catattgctg attaaaaaca gtgtagtgag agctcagtaa
                                                                    1560
atgittattg aatagataaa tccatggtig tagicatgat cattgacata atatgctccc
                                                                    1620
tttaggaagg tggatatcta aaaatgtgtg aatcaggtgg aatgttttgt cacatgctca
                                                                    1680
ctgctttcta ctctagatta ccgcttaaac cccatgttaa gaaaagcctg taaagctgac
                                                                    1740
attectaaat tetgteacgg tateetgact aaggeeaagg atgatteaga attagaagga
                                                                    1800
caagtcatct citgcctgaa gctgagatat gctgaccagc gcctgtcttc agactgtgaa
                                                                    1860
gaccagatec gaatcattat ccaggagtec gecetggact accgcetgga tectcagete
                                                                    1920
cagctgcact gctcagacga ggtgggattt gcgtgcaaaa ctggttacgc acagagctgc
                                                                    1980
tcagagaagt ttccactgga gaaaagttgt ttactttctc tcccttcagc cgtgaatgat
                                                                    2040
ctggtgaatt gaaggecate ttetaggete tecatggtet geatteetgt tetttgtaac
                                                                    2100
actgaattca acttggcatt agtcctgaca ctctaaagcg ttgttccata tttctctgtt
                                                                    2160
gaacaagggt gitcitical tatagetete tgtaaattig ticiteeeti ettettatte
                                                                    2220
tggatggtaa acceaagace tgceagaaag ataaaagtge ttteagetgg geaeggtgge
                                                                    2280
teacgeetgt aateecaaca etttgggagg eeaaggaggg tggateatet gaggteagga
gttcaagacc agcctggcta acatggagaa atctgtctct actaaaaaata caaaaaatta
                                                                    2340
                                                                    2400
gecaggegtg gtggegtgea ecagtaatet eagetaetea ggaggetgag geaggagaat
                                                                    2460
cacttgaacc cgggaggcgg tggttgcagt gagctgagat catgccactg caccccagcc
                                                                    2486
tgggcgacag aggaagactc tgtctc
```

<211> 2568

<212> DNA

<213> Homo sapiens

60	ctgctcacac	tcacacacac	acacatacac	acacacgctc	gctcacccac	acatacacat
120	atacacatgc	atgctcaccc	tcacacacac	atacacatgc	atgctcaccc	atacccatac
180	gtttacccat	gctcatacac	acacacacat	acacatgete	gctcacccat	tcacacacat
240	cacctgctca	cgattacatg	cccacataca	acatacacct	acacacgatt	acacatgctc
300	atgctcacaa	atacacatac	atgctcacac	ttacatacac	tgctcacaaa	cacacacaca
360	gcctactcac	aattacatac	atgctcacac	tcacacacac	atacacctac	acatgattac
420	ctcacacgat	cacacacatg	tgctcataca	tacatacacc	tcacacaaat	acacacatgc
480	cgagtacata	cacatacaca	tacacatgct	tgctcaccta	tgctcccaca	tacatacaca
540	cattcacatt	tgctcacaca	tacatacaca	cacacacgat	cacatgctca	cacatgctga
600	tacatacaca	tacacacgat	cacccataca	ctcacatgct	cgcacacatg	cacatgttca
660	tgctcttacc	ctcatacaca	ctcacacagg	catacacatg	tacacgatta	tgctcataca
720	caggcacaca	catgatcaca	cacaagcaca	catgctcaca	cacacacaca	catacatgct
780	ctcacacagg	ctcatacatg	tacacacata	tgcacaagtt	tacacaggct	tgatcacaca
840	caggattaca	taggcacaca	ctcacatgta	tgcacacact	tgctcacata	catacacaca
900	tttatacaca	tgttcacacg	cccatgcaca	tacacataca	catgtgctta	catgctcaca
960	tctgtctctc	tgccacatgc	acacacacgc	ccatatactc	acacacaccc	catatcacac
1020	tcggaggctg	ggtcttccct	tttccctcgg	tgctgtcccg	gacagttccg	acacatacct
1080	gggcttttct	caaggaggta	actgagcagg	ccttgaaggc	gagcatccag	cagctcgtct
1140	ggaagggcag	ccaagggcag	ctcatcccca	gtacggcttc	ctgcgttgca	ctggggaggc
1200	tcggccttgt	ctgtctcgtg	ctgggagcgt	tacagaacac	cttacctgaa	gagtctagga
1260	teceteagaa	tgttggcccc	ctaagccact	gacttgcgcc	tgcagaccag	gctgagcatg
1320	tgccataaag	cggtaagaaa	acaagggagg	tcacccacag	tggcttggcc	tcatgtgcag
1380	cctttccaat	tccacacttg	accigtgcag	agcaccttac	cctgaaggac	agacgctgag
1440	gagaaggcgc	cccagaagcg	gacaccaagg	cacgtccctt	atttcatcct	tegeegttge
1500	ctgccccgtg	tetettgtta	ttgcttttca	ggaagggagt	atgctgagtt	acgtcgcata
1560	ctggctgctc	ctgcatggaa	aggagagaac	attttgtaaa	caagtgacac	aacaactttc
1620	ctcctaactg	actgccagga	atgctgagtt	aggctgcttc	tccctgcccg	tggggctgtc
1680	atcctcccag	gagcacctac	gtggctacct	caggattacg	gcgctttctc	tcttctctct
1740	ccaataacag	tgctctctct	cctgcggctc	cagacattca	aaatcaaggc	caaagggaga
1800	ggcaccaggg	gggcetccat	agaggtactt	gcgttttccg	ctatgcctct	acttcaaact
1860	acgccacggc	agagaccggc	ccggcgcccg	gcgcctcctc	ccccgcagca	acaacgctgt
1920	gcccacagga	ccccacgcga	tggccgcagg	caaaggatgg	aggaacagag	ccctccccca
1980	actageetet	accegeceae	gaagcatgag	ccaccaagcg	tagattccag	caccggcccc
2040	cgcagcagtt	gtgccatgga	tcacgtcgca	acaccetgte	tagggacatc	gtgttcccgt
2100	ctglaatgag	ttgctacttc	atatatgggc	gtgcgtgttc	tgttttagca	acagcaccat
2160	cttactgtag	tgtacagagt	aacgcttttc	gctacaagaa	atggtgaggg	gacgttcaac

ctacgctaat	ggttaacctg	atagaattaa	ctcgtatttt	tctatggttt	taacctgatg	2220
ctccactgtc	tccgtcatgg	ggttgttttg	ctgtttgggg	ttgggccttg	tttccctttc	2280
ctttctccag	tccacgtgta	gactttgcgc	ttgtttggat	gaagaagcag	atcggaagta	2340
actgctccct	cctcaaggtt	gtcttcagac	gtcttggaga	cgttcctaaa	cactgagggg	2400
gaagacagcc	aatagcaccc	attaaaagaa	atacctaaat	aaaacctctc	tcccactcag	2460
ctatgctagg	gcttggctgt	aggtgtgcac	tgtctattta	catccgtcct	tacaaccatc	2520
cttgtcctcc	ttggtaccgt	atcaagctct	ttcccatgac	atttggtt		2568

<211> 2425

<212> DNA

<213> Homo sapiens

gaaacagtag	aaaaggccga	ggctccatta	ataactgaga	gtgcttttga	tgctggtttt	60
gagaaacttc	ttaaagaaat	aactgaagct	cctccttatc	agccccaggt	gtcagtgaga	120
gaagaaactc	acgagaagga	gtcctcacag	tcagagcaga	ccaggttctt	ggggacagtg	180
ccccattttt	acagggcagc	ctcacagacc	tctgaaatga	aggataaaag	taatggtttg	240
gaatctcaag	tcaaccaatg	tgataaaatg	ttgggaggag	acgcacttgt	gactgattta	300
ttggtagatt	tttgtggttc	cagaagtgga	gttgagatcc	ctagaacccc	acaactttat	360
gtggctcatg	aaatagggac	cattaaaact	gtaacccccc	cagaggacag	ggacagtgaa	420
agtggggttg	tagggggaca	agggactctt	caggaacctg	gctttggaga	ggcttctgaa	480
gcaattagtg	tgtccagaaa	taggcaaccc	attcctctcc	tgatgaacaa	agaaaactct	540
acaaaaacaa	gtaaagttga	attgactcta	gcatcgccat	atatgaaaca	agagaaagag	600
gaagaaaaag	aaggtttctc	tgagtctgat	ttttcagatg	gaaacaccag	ttctaatgca	660
gagagctgga	gaaatccttc	cagttcagaa	gaagaaccca	gtcctgtttt	gaaaactitg	720
gaaaggagtg	ccgctaggaa	aatgccttcc	aaaagtctag	aagacatttc	atcagatica	780
tcaaatcaag	caaaagtaga	taatcagcca	gaagaattag	tgcgtagtgc	tgaagatgat	840
gagaaaccag	atcagaagcc	agttacaaat	gaatgcgtac	caagaatttc	cacagtgcct	900
acacaacctg	ataatccatt	ttctcaccct	gacaaactca	aaaggatgag	caagtctgtt	960
ccagcatttc	tccaagatga	ggtgagtggc	agtgtgatga	gtgtttatag	tggagacttt	1020
ggcaatctgg	aagttaaagg	aaatattcag	tttgcaattg	aatatgtgga	gtcactgaag	1080
gagttgcatg	tttttgtggc	ccagtgtaag	gacttagcag	cagtggatgt	aaaaaaacag	1140
cgttcagacc	catatgtaaa	ggcctatttg	ctaccagaca	aaggcaaaat	gggcaagaag	1200
aaaacactcg	tagtgaagaa	aaccttgaat	cctgtgtata	acgaaatact	gcggtataaa	1260

attgaaaaac	aaatcttaaa	gacacagaaa	ttgaacctgt	ccatttggca	tcgggataca	1320
tttaagcgca	atagtttcct	aggggaggtg	gaacttgatt	tggaaacatg	ggactgggat	1380
aacaaacaga	ataaacaatt	gagatggtac	cctctgaagc	ggaagacagc	accagttgcc	1440
cttgaagcag	aaaacagagg	tgaaatgaaa	ctagctctcc	agtatgtccc	agagccagtc	1500
cctggtaaaa	agcttcctac	aactggagaa	gtgcacatct	gggtgaagga	atgccttgat	1560
ctaccactgc	taaggggaag	tcatctaaat	tcttttgtta	aatgtaccat	ccttccagat	1620
acaagtagga	aaagtcgcca	gaagacaaga	gctgtaggga	aaaccaccaa	ccctatcttc	1680
aaccacacta	tggtgtatga	tgggttcagg	cctgaagatc	tgatggaagc	ctgtgtagag	1740
cttactgtct	gggaccatta	caaattaacc	aaccaatttt	tgggaggcct	tcgtattggc	1800
tttggaacag	gtaaaagtta	tgggactgaa	gtggactgga	tggactctac	ttcagaggaa	1860
gttgctctct	gggagaagat	ggtaaactcc	cccaatactt	ggattgaagc	aacactgcct	1920
ctcagaatgc	ttttgattgc	caagatttcc	aaatgagccc	aaattccact	ggctcctcca	1980
ctgaaaacta	ctaaaccggt	ggaatctgat	cttgaaaatc	tgagtaggtg	gacaaatatc	2040
ctcactttct	atctattgca	cctaaggaat	actacacage	atgtaaaagt	caatctgcat	2100
gtgcttcttt	gattacaagg	cccaagggat	ttaaatataa	caaaatgtgt	aatttgtgac	2160
tctaatatta	aataagatat	ttgaacaagc	taggaaaatt	gaatttctgc	tgctgcttca	2220
aagaaaaagc	tgccccagag	cattaaacat	ggggtattgt	taagaagcaa	aatgttcttg	2280
tttgccatca	tgtgtttcac	accacaattc	tgtgccacag	ttaagagggt	ctggtaccct	2340
tgcaggacct	ttgtaggttg	tgggaaaaaag	tegeagaaag	atactcaaag	tggagcaggg	2400
aatggagaca	gacatcagtg	atgat				2425

<211> 2194

<212> DNA

<213> Homo sapiens

gagggagtgt	ggaagtatcc	caggaacaga	gactgtccca	attggaggcc	ccagagaggg	60
aactggggct	tgtccttggg	gccactgcta	ctgctggggc	ttggggagga	ggcctgaggg	120
caaccctgaa	gccgtggagc	agatggcctg	tgtggcagtg	gctgtcagat	tggagggcag	180
aatggaatct	gatgtggctc	cagcttcttt	gaaggctgag	tgcacatctg	gccagctgct	240
ggtgatgggg	aagggggtca	ctgctctatt	gagctatgtg	tgactcaccc	ttttccctag	300
tatcctgtcc	ttgtgctatg	tcaggaaacc	tectecceag	ccccaccgag	tcttcaaagg	360
agttggtccc	ccaggagctt	ttgtcctggg	gatttcccat	ccgcactggg	agcactgtgg	420
gtagatgggc	attccaagtt	cctgcccaga	cagaagccat	gggcatggag	acctggacct	480

ttcagcttgg	aaaagggaga	ctgaggctga	gtggacacag	gacacacaca	cactctcagg	540
gtcacccttg	gggcagcaaa	gtggagcatc	ttggctcccc	atttaatgag	tgacccaact	600
ggtgactcat	taacatccca	gagcacaatg	aaaggcatgg	gggcggggca	gccctccggc	660
tctgtccctc	atcccttgcc	caggcatctg	ggtttcctgc	agattctcag	ctcagctgct	720
gctttttctc	actggaaaag	ggctcccagg	gctatgtggg	atcagggctg	accactctat	780
aaactgatga	ggaccccagg	tctggccaag	tgaccctgtt	ttaccttcgt	tcaggaagag	840
ctgaggccta	ctgtgtacta	ggcacgggag	atatagtggc	aaccagattg	gctagctttt	900
ctgtcctgga	ggccacagcc	tacttaggag	cagaaatagt	tatgaaataa	acacacaaat	960
aagtggaagt	gttggctggc	cgatgcagga	aatcaataag	gtggtctggt	ggtgacttgc	1020
tgggaatgct	tctttgagta	gggaggtaac	agaggcctct	gagaggctgg	cagctctgag	1080
tcaagtctgg	gattcctgga	ctgtgggctc	tettgccccc	atctccaggg	gacttgtggc	1140
agccagagag	ctgagggtgg	ttgttgggga	ggaaggcctt	gggcttcaag	accaccctgt	1200
ctgtgcaatc	ctcacaggcc	caccgtggtg	cacgcaaacc	acttcctggc	catgcgctcc	1260
ctcctgcttc	tcagcgcctt	ctgcctcctg	gaggcggccc	tggccgccga	ggtgaagaaa	1320
cctgcagccg	cagcagctcc	tggcactgcg	gagaagttga	gccccaaggc	ggccacgctt	1380
gccgagcgca	gcgccggcct	ggccttcagc	ttgtaccagg	ccatggccaa	ggaccaggca	1440
gtggagaaca	tcctggtgtc	acccgtggtg	gtggcctcgt	cgctggggct	cgtgtcgctg	1500
ggcggcaagg	cgaccacggc	gtcgcaggcc	aaggcagtgc	tgagcgccga	gcagctgcgc	1560
gacgaggagg	tgcacgccgg	cctgggcgag	ctgctgcgct	cactcagcaa	ctcgacggcg	1620
cgcaacgtga	cctggaagct	gggcagccga	ctgtacggac	ccagctcagt	gagcttcgct	1680
gatgacttcg	tgcgcagcag	caagcagcac	tacaactgcg	agcactccaa	gatcaacttc	1740
cgcgacaagc	gcagcgcgct	gcagtccatc	aacgagtggg	ccgcgcagac	caccgacggc	1800
aagctgcccg	aggtcaccaa	ggacgtggag	cgcacggacg	gcgccctgct	agtcaacgcc	1860
atgitcitca	agcgtgagtc	gggggcgcgt	tcaggggtcc	tectectect	cccaggaccc	1920
cctgcaagag	ttaggacgac	attccgtgcg	ctccattctt	cactgcctct	catttatgct	1980
gtgacaaccc	agggaggcag	gactgtcact	cagctttttg	tacagactgg	aaactagatt	2040
cagagacagg	tagcagcgtg	taaaggaatg	gttcagggag	cggaggcccc	agagggactc	2100
catggaatgt	attccgaccg	aatttcgtca	aagtgctcgt	ccttgtgtat	ctiggaatga	2160
ataacattta	ataatccatt	ctgcctcagt	agtg			2194

<210> 4674

<211> 2876

<212> DNA

<213≻ Homo sapiens

(100) 1011						
aaagggccag	cggggcacgt	ggctcgggac	gcagttcgct	gccgcccggc	agtagctctc	60
aggttaggcg	ggtcccgctc	cgcttccgcc	gtcgctgccg	cgccgccccg	ggcccgacag	120
gccgggtcca	gggactgcaa	cccagcgagg	gacgcgggca	gccatggccg	aagcggcgcc	180
tgcccgggac	cccgagacag	acaagcacac	agaggaccag	agtccttcga	caccettgee	240
ccagccagct	gctgagaaga	actcgtacct	ctactccacg	gaaatcacac	tgtggacggt	300
ggtggccgcc	attcaggcct	tggagaagaa	ggtggattct	tgcctgaccc	gcttgctgac	360
tctggagggg	cgcacgggga	cagccgagaa	gaagctggcc	gactgcgaga	agacagctgt	420
ggagttcggg	aaccagctgg	agggcaagtg	ggccgtgctg	gggaccttgc	tgcaggagta	480
cgggctgctg	cagaggcggc	tggagaatgt	ggagaacttg	ctgcgcaaca	ggaacttctg	540
gatcttgcgg	ctgccccgg	gcagcaaggg	ggaggccccc	aaggtgcccg	tgaccttcga	600
tgatgtggcc	gtgtatttct	ctgagctgga	gtggggcaag	ctggaggact	ggcagaagga	660
gctctacaag	cacgtgatga	ggggcaacta	cgagacgctg	gtctccctgg	attatgcaat	720
ctccaaacca	gacatectea	cccggataga	gaggggagag	gagccttgtc	ttgaccggtg	780
gggccaggag	aaggggaatg	aagtagaggt	gggacgtcca	aggatgatgg	gcactggcct	840
ccctccgtat	ccagagcacc	tcaccagccc	acttagccct	gcccaggagg	agctgaaaga	900
agggcaggcc	cccaagcagc	agcaggactc	agaggcgaga	gtggccccag	ccgggccaga	960
agcaggactg	gcattgcgga	ctgacctcca	gggagaggcc	cagatctgat	ctgcagagat	1020
ctcccttccg	gctatcgtgg	ctgctgttca	ggtggtggag	aagaagatgg	aaccccaggc	1080
tgcctggcta	cagageetgg	agggatgcat	gtggacagcc	aagaagaagc	tggctgactg	1140
cgagaaggtg	gctgtggagt	tcgggaacca	gctggaggcc	agcccaccac	caagccagat	1200
ctggcaccag	tggagaaagg	agacaggcag	gcaactgcag	cctgcccagt	gcggagaggg	1260
gccagagccc	gggcactgat	ttctccagga	gagagctttg	tttaccccca	aggactgggc	1320
ttcactccta	aaagctgctt	ctaaaccctt	tgcgtgcagg	ccccagacag	tgtgggtgca	1380
tgactgcttt	atctgttgga	tgtttccaca	ggcaggctgt	tgctgccaga	attclagaag	1440
cagtgtaaac	gggctgctca	ttcattaggg	ccctggttaa	tggtggagcc	agcctggtgt	1500
gtggcagcca	ggtcgggctg	gtgtcttcag	agtggctgtg	tcagaagggc	ctcgtgtatg	1560
acagaaacgc	aggccagctg	gaacgtcccc	aggcgttctg	tcctggatca	agcaagagga	1620
aggggcaagt	gggaggagcc	agcagggccc	acaggagacc	actgtggcac	acctgtgctc	1680
aggcagcatg	gtgctgggcg	ctgcaggggc	tcallaatgc	atgagacaca	gattccactc	1740
aggagagaga	agacgaacac	agacacagta	cacggtggga	aggggcactt	ccatgicige	1800
agccagagag	ccctttcaga	atctgaccca	aaagatggcc	agtattctaa	gaaaaagcac	1860
atatgcttac	gatgtgcaaa	cagctgaagg	tgttctttca	actatagtaa	aaggggtgtc	1920
teggtggcca	caggcggccc	agatgaagag	acctgtctgg	gccgcagaga	ggagtctgga	1980
ggccccccga	gccaggctga	gccagccgct	aggggcacgg	agcagtgccc	accttgcgcc	2040
cagtgtggcc	agagcttcgg	ccggaaggag	ctcagtgcgc	cgcaccagcg	cgtgcatcgt	2100

ggcccccggc	ctttcgctgg	tgctcagtgt	cccaagagct	tcacgcagcg	gaccaccccc	2160
gccagccacc	gtcgggcgca	cgtggccgag	tgcacctaca	cccgcgccca	atgcggcaag	2220
accttcctcc	agcagtcgac	gctcacaccc	actactgcgc	gcacatcagg	gagaagccct	2280
acgagtgcgc	caagctcttt	ggccgcctgt	ccacgctgct	ggagcaccgg	cacatgcaca	2340
cgggtgagcg	gcccttccag	tgcacgcaat	gtggctgctg	cttcagccgc	ctgtccacgc	2400
tgctggagca	ccggcacaca	cacatcagcg	agaagccctt	ccagtgtgcg	cagtgcgaca	2460
agcgtttcac	gcgtctggcc	aacctgaccg	tgcaccagag	agtgcacttg	ggcgagcgct	2520
ccttccagtg	cgcccagtac	agcaggagat	tcatgcagaa	gcccggcttc	ctgcgtcatc	2580
tgtgtggcca	ctcgcaggag	aagcactatc	cttgcagccc	ttgtggttcc	catgatcagt	2640
ggtgaaatat	agtgattttc	acctgtgctt	ccattctgaa	gttctggaaa	gaagtactgg	2700
atggactgaa	gtccaggaca	acgttccaaa	gaaaggcaga	gtccaggtag	gcttggagga	2760
ccaagccctg	gatgagcact	ggagggcaga	ggcctcagtg	tccagcactg	tgccctgcac	2820
atggaaagcc	cctacgtttg	tggaatgaat	gaataataaa	aatgttttca	taagtg	2876

<211> 2262

<212> DNA

<213> Homo sapiens

attcatcagc	agtttccgtc	agctcttcct	gcaaaacaca	ttcaagcccg	accatatgtc	60
accaccccca	ctgctattcc	caggtcccag	ccaccatcat	gtctcacctg	gactattgca	120
cgcacctcct	acttggtctc	cccacttctg	cccggccgg	ggctggtgtg	ttctccaccc	180
agcaacacag	cagtcagagg	gatcctctta	caacataagt	cagatcacct	cactctttgc	240
tcaaaatcct	gcaatggctc	ccatctcctt	cagagtaaat	tccaaagccc	catatctgcc	300
cactgccccc	cactctagaa	ccttccctcc	tgttgttctt	ccctggttc	cctccattcc	360
agctacaggg	atctccttgc	tggctcctga	gcttaccagg	tatggtgctg	cctcagggcc	420
tttgcacttg	cagtgactgc	tgcccagtaa	gtttggtaag	gttggtcccc	agacatccac	480
accgccctct	tctgcacctc	ctgctcagat	gtcactttct	caatgtggct	tttgctattt	540
aaaactgcac	ccctttaaat	tgtacccagt	ttaaaactgt	accecctace	ctaccctcct	600
gatggagttt	ggatatttgt	ccctccaaa	tctcatgttg	aaatgtgatc	cctgtcaggt	660
gtcatggttc	acgcctgtaa	ttccagcact	ttgggagacc	gaggcaggca	gatcacctga	720
ggttaggagt	tcaagaccag	cttggccaac	agggtgaaac	tccatctcta	ctaaaaatac	780
aaaaattagc	tgggtgtagt	ggcaggcgcc	tgtgatccca	gctactcagg	aggctgaggc	840
aggaggattg	cttgaactcg	ggaggtggag	gttgcagtga	gccgagatcg	cgccactgca	900

ctccaggctg	ggagacaaga	gcaaaaactc	tgtctcaaaa	aaaaaaaaaa	aaaaagcaag	960
aaccagccca	gggctgagtt	atgaagaggg	gtgctgcaga	ctaggccagg	tccccaacct	1020
tgtcaggtct	gcaatgccaa	gcttccagac	ctgatcacaa	aggagtggga	ctccaagaag	1080
agcgcgatgg	gatgtctggg	tgaatgcaca	gtctccacct	gtggttcccc	agatttacct	1140
ggaccctctg	ggcctgcaga	attgcccaag	tcctccctca	gaggataaca	ggctctctcc	1200
ttgcctaaag	atgataccgc	ggccttcaac	gtggaagacc	aaacatcctg	ccctccgaat	1260
ccacctataa	cctgtgccgc	ccaccagatt	catcactggg	gttaagtcac	aaccacaccc	1320
acctggggac	atgctgggcc	agccagggag	gacaagacta	caccccaaag	gagctatgga	1380
cccagccagc	acggagcagt	gggggtggga	cagtgctctt	gggaccggat	caagggcagg	1440
aggagctcat	gaaatggggg	acttctccca	ttgttatgga	ctcaattgta	ttccccaaaa	1500
aaattcatag	atggaagctc	taacctccaa	tgtgactgta	tttggagata	gggcctataa	1560
ggacgtgatt	aaggtgaaat	gaggtcatca	gggtgagacc	ctaacctgat	aaggctgtta	1620
tacttagaaa	acaaggaaaa	gacaccagat	ctcattctct	ctctctctct	ctccctccct	1680
ccctcttcct	tccctcccc	ttcctcctgc	cttacatgca	cagaggaaag	gctatatgag	1740
gacacaggga	gaagagagcc	atctacaagc	caagaagcga	gccctctcca	gaaaccgatc	1800
ctgctggtac	cttgatcttg	gacttccagc	ctccagaacc	aggagaaaat	aaatgtctgc	1860
tgtttaagcc	gctcagtctg	tggtttcttc	ttatggcagc	ccgagatgac	taacacacca	1920
gtcatgccgg	atttaacagc	aggagacagt	gctaacgcac	tgctgaagtg	gctcttgaaa	1980
cttggagaaa	gatggcagat	gacagaagaa	gggatcaaaa	gacagagaag	tgggcatgga	2040
ccagagcagg	cagaaaaaccc	actgggtgac	agcgttccac	aagacaccac	caaggacgtc	2100
ccactgaccc	agcaccggct	tcaccaggta	gctcagggat	ggggctgtcc	gccataggcc	2160
agggtgggtg	gtgggaggtg	ccgttacaga	accaggtttg	ctgagagagc	tgggaataag	2220
agagagagag	aaagaaagaa	agaaagagag	aaagaaagaa	ag		2262

<211> 4486

<212> DNA

<213> Homo sapiens

ataaaaataa	aaaaaaataa	cagccacaaa	ctagggagaa	cgaaaaaacag	acacttccca	60
		_				120
caagagggca	rggggarggc	cagiicaagc	catcaggete	attagcaatt	agagaaatgc	
aattaaacca	caatttaaaa	gccaaaagct	ccacaccctt	agaatggccg	aaattaaaaa	180
gatggacgga	cgccagcaag	tgttgggaaa	gacgcggagc	ccgctgcaca	tgagggtgtg	240
gggggcacca	tggcccctgc	cgccgtctct	ccagctgccc	tccctccctg	cctccctccc	300

tctgtgtcca	cctgctcctc	ttccagttcc	ctctgcctgg	ctgtttcctc	tagtgaatgc	360
tcactcagct	ggcaggtcct	ggcttagcaa	agctcacagg	cattcggctc	tgcatctgcc	420
tgcggctctg	accctgtctt	tgtggccctc	tcaccagcca	gctgccccc	ttgactctga	480
gctcccagag	ggcagggaag	cctctgcctg	ggcacgtctt	tgtacgtaca	gcacctaaca	540
tgggcctggc	gcttaggagg	gcccaggagg	cacctgagta	agggcaccag	tcacctgctg	600
ctacagcaga	agtgtttcct	cctaacagtt	ttgtgctgct	ctcctctta	gggcatcttg	660
acccagccga	aaaagttgaa	gatgctcacc	ccaagttatg	gtgtgctctg	agcgaaggca	720
aggtgaccgt	gttcaatgct	tcttcatgga	ccatccacca	gcactccttt	aaagtgggca	780
ctgcaaaagt	gaactgcatg	gtgatggccg	accagaacca	ggtgtgggtt	ggctcggaag	840
actccgtcat	ctacatcatc	aacgtccaca	gcatgtcctg	caacaagcag	ctcacagccc	900
actgctccag	tgtcacggat	ttgattgtgc	aggacggaca	ggaggcaccc	agcaacgtgt	960
actcgtgcag	catggacggc	atggtgctgg	tgtggaatgt	gagcacactg	caggtgacca	1020
gccgcttcca	gctgccgcga	ggtggcctga	cgtccatcag	actgcacggc	ggccgcctgt	1080
ggtgctgcac	aggtaacagc	atcatggtca	tgaaaatgaa	tggatccctc	catcaagaat	1140
tgaagattga	ggagaacttc	aaagacacca	gtacctcctt	cctggccttc	cagctccttc	1200
ctgaggagga	gcagctgtgg	gcggcctgtg	caggacgcag	cgaggtttac	atctggagcc	1260
tgaaggacct	ggcccagccc	ccgcagaggg	tgcccctcga	ggactgctct	gagatcaact	1320
gcatgatccg	ggtgaagaag	caggtagggt	ggagggcccg	ccatccccag	catccccggc	1380
aggtctcgct	tgccctggcc	gcatccccat	gctcccgaga	gccagccgca	cgccctcgtg	1440
ccctcctgcc	ttcgcctctc	agggttcccc	tcctaacagg	cacgtgcagg	gttggtgcca	1500
acgggcccac	aggtcaccat	gtgctggttt	catggtgtgg	tecaageact	ccacagcgca	1560
ttcccccaaa	ctggatggct	tcaaatgaca	gaacttaatt	ttctcacagt	tctcgacgct	1620
ggaggtccaa	aaccaaggtg	teggeagage	cgcttccctc	gggggctcgg	ggggagcgtg	1680
cactccaggc	ccttctctcc	cccacccgct	catgtggccg	ccgtccttgg	cgtgccctgg	1740
ctgtggacgt	ggcactctaa	cctgtgcctg	caccacatgg	cactctctct	ccctgtgtgt	1800
gtctcttatg	gggatgctgg	tcgtactgga	ttaaccagtg	tgaccacatc	tgaactatct	1860
gcagactcta	cttccaaaga	agctcacatt	ctgaggttca	gaggatatgg	atttgggggg	1920
acactgttca	acccattaca	gtctgccctc	tggcccccta	aatccacatc	tgttccacat	1980
gcaaaataca	cccctccat	cactacatcc	cccgaagtcc	cagtecetee	agtgtcagtt	2040
ctgagcccca	aatctccaag	tgaagccaaa	gccctccgat	gtgggtgagc	cgtgggcggg	2100
teccacetgg	ggcagaacca	acccctcttg	caccgtgaac	ctgcgcagct	agaacaggct	2160
gtctcttccc	aaaacacaag	gctgggccag	gcagaggtta	gacattctca	tccgaaaagg	2220
gagaaattgg	aaggaaagct	ggacttccag	gccccgagca	agteggagae	acggcggggg	2280
tggggcggaa	ttcgtttagt	gttttcccc	ataacttaat	tggggtataa	ttgatgaata	2340
aaaattatat	atatctaagg	tatacaaggt	gatatttcag	tatacattgt	gaaataatca	2400

caagctaatt	accatatctg	tcacctcaca	caggtggctg	ttgtttggtg	agaacactta	2460
agatgcaccc	cagcactttt	ccagcacgca	gtgcagtctt	cttccccaga	ggcccctcgg	2520
ccattaggtt	ctaagcccga	gaagggtcct	gtgcacctgc	caggcggccc	tgaccctcag	2580
gcggcagagg	ccctgtccag	gaaggacata	gatttggagg	cacacgcgtc	aattctgggt	2640
agccgggcaa	agccctggtg	ctcttgggag	gggcttcttt	ccctgctgac	ccctcccgtt	2700
tctccctggc	aggtctgggt	gggcagccga	gggctggggc	agggaacacc	caaggggaaa	2760
atctacgtga	ttgacgccga	gaggaagacc	gtggagaagg	agctggtggc	gcacatggac	2820
accgtgagga	cgctgtgctc	ggctgaggac	agatacgtgc	tgagtgggtc	gggcagggag	2880
gaggggaaag	tcgccatttg	gaaaggcgaa	taaacgtggc	tgagtctgcc	aagtggaact	2940
gtgccctatg	tgtggggact	ggctgccccc	tagagcctgc	caggagcaga	agcctggagg	3000
ggtggcaggg	cagagcagcc	caggctcagc	atggagccca	cttaccgtgt	ggccagccgc	3060
gagacccatg	gccacgcacc	ttctctcagg	ccttcgggcc	ccctggttaa	actgcaccaa	3120
gggtgtttcc	tgttggggtg	tgtctcaggc	aggcagctgc	gtcttgttgg	tgataacctc	3180
tgctgggagg	ttactttgtt	gcctagaaag	ttctggaatc	cacaaccagg	ggctggcact	3240
ggagccagca	gcttggccga	gtcacaggtg	acccgtggcc	ctcacgtctc	tggttttacc	3300
tttccttact	tcattcattc	actcacccag	tccttacgaa	tcaccgagga	acactgggct	3360
gagcacatga	cagggagcct	ggagccccgg	ggcctccagc	gaggcctgag	aagggtggtt	3420
cgggtaacca	ctgtgggctc	tctcccatca	cagaaggtgg	acagggccta	cccaggtgga	3480
ggggaccacc	ctgcgatcag	gtgtttgcga	caggggttgg	gccagctgag	gcaagctgtc	3540
ttttttccct	tttcttttta	atagatgcaa	catttttata	ataatcctag	agaccttttt	3600
tctaccaaag	atcacagacc	agaaaaagtt	ccatctaaaa	tatcatgccc	aggaaagcac	3660
atgggatcaa	aagtaaaata	gcatcatgtg	tgatctcgtc	ttccagcgtg	ccgctcagtt	3720
ccccgaatcc	gtgtgcacac	gtgtgatctc	gtcttcagtg	tgccgctcag	ttccctgaat	3780
ccgtgtgcac	actgcgtatg	tgtacgcgca	gcatgctata	ctgaactcaa	caagatcttg	3840
gctgtacata	atatttgtaa	aagagaccct	ttgcaccitt	ttactgtaat	gttgagactt	3900
cattacttaa	atgttctacg	gaaggttctg	gtgtggttgt	tggagccgga	gggagcgtgt	3960
cagcacgtgc	tgagggcatg	gggcctgccc	cctgggcacc	catccacaag	ctgggccacg	4020
gagctccagc	ttctcaggac	aaagccccgg	ggctggcgca	tcctgagggt	ctctgggggt	4080
gtitgccagg	ctcctgggat	gggccgcttt	cagaagccct	gcagtgcctc	cagatggaaa	4140
ggcgggcccg	gcctccggtt	gggtctgcat	tttggagagt	ccacaccacg	gaccaggttt	4200
teccecaagg	cttggctttg	tgtagctact	aacttcttgg	ggcattctga	gagtgtgggc	4260
agagagaatt	atgtggcctc	atcctcccc	aaggctgtgc	ttgcagcccg	ggcaccttcc	4320
cactitciag	ctctggagag	gttggatttt	gcttttgtaa	acacatgaat	ccttatgata	4380
aaagtetgte	agtcaaaaat	acatttataa	attatttaat	gccagtcctc	atgtaacctc	4440
aggtatette	agcttgtgga	gaataaatct	ggtttaataa	acactg		4486

<210> 4677 <211> 3453 <212> DNA <213> Homo sapiens

agcaccttgc	tggcccacct	ggactgagaa	gcatcagagt	cctggtaggg	gtaggcccag	60
tagagcgcat	ggcccccgtg	accctcacag	ccagagttgg	cgcggtgggt	gggaattgta	120
ggtagatttc	agccagcttt	agggagcagc	tgttttaggg	tcactgctgc	ccagccatgg	180
agtgagctct	gtgtcactgg	gggtatgcaa	gcagggacag	gagacgtact	tgtcaaggat	240
atggtagcag	gaagtctggg	ttagactgga	agacctcctt	tctgtgcgga	tctgattcca	300
gcaggaagcc	acggccaggt	ggccgagggc	catgctgggc	acccaagcag	ccctgactgg	360
tgctgtgggc	gggcaggccc	tcaacatatc	cttcctggaa	gaggacatta	acgctcgtga	420
tgtttccaca	cctaccctca	cgggacccca	gggagtggcg	gctgggtggg	gaggctcacg	480
acagcatcat	ccttcccaca	ttctgagtgg	cccctgagcc	ctgattcttc	ttttccagcg	540
ataagttcac	acggttttgc	cagtggaaga	atgtggagct	caacatccac	gtgagtgggc	600
ttgggtgggg	catggaaagc	cacgcaccct	gctgctcctc	tcccgggagc	tgggcctgtg	660
gcttggctgg	gagggggagg	tcaggggatg	tctgtccttt	agcccccagg	gccgtggcta	720
tgggggtcag	ggccgggatc	ccagcatggg	gaggccggag	caggtaaata	tgtggcaagg	780
atggccagga	catgggtatg	gggaccctgg	catggggcca	gccctgctg	cccaggtgcc	840
tctgccccag	ggctgggcag	aggcagcctg	tggtgaccgc	agctgtcgct	gcccctcagc	900
tgaccatgaa	tgacttcagc	gtgcatcgca	tcattgggcg	cgggggcttt	ggcgaggtct	960
atgggtgccg	gaaggctgac	acaggcaaga	tgtacgccat	gaagtgcctg	gacaaaaagc	1020
gcatcaagat	gaagcagggg	gagaccctgg	ccctgaacga	gcgcatcatg	ctctcgctcg	1080
tcagcactgg	ggactgccca	ttcattgtct	gcatgtcata	cgcgttccac	acgccagaca	1140
agctcagctt	catcctggac	ctcatgaacg	gtggggacct	gcactaccac	ctctcccagc	1200
acggggtctt	ctcagaggct	gacatgcgct	tctatgcggc	cgagatcatc	ctgggcctgg	1260
agcacatgca	caaccgcttc	gtggtctacc	gggacctgaa	ggtgagcgcc	cctgctgtcc	1320
ccaggctgga	cctccgtggc	tgtcctctcc	ttcctctcga	cateceggee	accaggecea	1380
gaggagtggg	gctcctggga	catggccgcc	ccgtatcttc	ccatctccgc	ccctgccctt	1440
cccaccgage	cactctctgg	gtccaggttg	tagctgggga	caggagagag	gaccccacc	1500
tttgcccttt	ctttgggtac	ccatcgtcct	ctccagtgaa	gcagtgaccc	agctggcatc	1560
ttgcctggct	gggccccatc	ctgagctgcc	ccaggcagct	cactgggctt	ccttcacagc	1620
cagccaacat	ccttctggac	gagcatggcc	acgtgcggat	ctcggacctg	ggcctggcct	1680
gtgacttctc	caagaagaag	ccccatgcca	gcgtgtgagt	gececcace	ctctccctcc	1740

ccaccccttg	ccactcccgc	ttatggcccc	cttgctccca	caggggcacc	cgcgggtaca	1800
tggctccgga	ggtcctgcag	aagggcgtgg	cctacgacag	cagtgccgac	tggttctctc	1860
tggggtgcat	gctcttcaag	ttgctgcggg	ggcacagccc	tttccggcag	cacaagacca	1920
aagacaagca	tgagatcgac	cgcatgacgc	tgacgatggc	cgtggagctg	cccgactcct	1980
tctccctga	actacgctcc	ctgctggagg	ggttgctgca	gagggatgtc	aaccggagat	2040
tgggctgcct	gggccgaggg	tgagtaccct	ggcgccttgg	gcatgctgct	ggctgtgccc	2100
ccatgaggac	aagggctgtg	tcccgtcacc	tggaaccccc	tccaaggtcc	cagcctcctt	2160
ggaggagctc	ataagttggg	gcatggccag	ccctgtccaa	tgttctcaga	gtggaggggc	2220
tgccctggga	gttggagctg	ctggaaccag	ctagtaactg	gcttccagag	gggcccttgt	2280
cacagatgat	gatgatagca	gctttttatg	tttatcaaac	atttattaag	cactgggcat	2340
agggttgcaa	aatttaccag	tgttaaaacc	cacaacccta	cgatgtagga	acttctgttc	2400
tcccatttga	gagatgagga	aacagctcag	gcttaaggaa	cttgcccaag	ttcacagagc	2460
cagaaagtgg	ctcaatcagg	agttaagccc	gggactcagg	gtggggctga	gcccagatga	2520
ctggcctctc	cccacagggc	tcaggaggtg	aaagagagcc	cctttttccg	ctccctggac	2580
tggcagatgg	tcttcttgca	gaaggtaaca	gtctgcggca	gggactgggg	gtgctctgca	2640
gccccacccc	cgagctaatg	cccatgaccc	ctgttctgct	gcagtaccct	ccccgctga	2700
tcccccacg	aggggaggtg	aacgcggccg	acgccttcga	cattggctcc	ttcgatgagg	2760
aggacacaaa	aggaatcaag	gtactgggcc	ttgcctggcc	tcttgtacct	aggctgtgat	2820
cctggcctgg	ggaaggatcc	ctctcccttc	ttatcacctg	tgagaccctg	tgccagccct	2880
gccagcttgt	aggcctcagt	tcctccttgg	ccaacttccc	tggggggtgg	cagttgcact	2940
gaccatccct	acccagggcc	ctgggtctgg	gcagcctgtg	gctgatggta	ttccggcatc	3000
tetgtccace	catgtgcccc	tgccccatcc	acctggtaaa	gttactggac	agtgatcagg	3060
agctctaccg	taacttcccc	ctcaccatct	cggagcggtg	gcagcaggag	gtggcagaga	3120
ctgtcttcga	caccatcaac	gctgagacag	accggctgga	ggctcgcaag	aaagccaaga	3180
acaagcagct	gggccatgag	gaaggtgagg	gtcgccggct	gctgcggcac	caggcccctg	3240
cctgcttaga	agtgagcagc	tggctcgggt	ttaaggaact	caccctggat	cacagccaga	3300
aagtggcggc	tetgggatte	aaactcaggc	tggggccggg	catggtggct	cacgcctgta	3360
atcccagcac	tttgggaggc	caaggagcgc	agatcacttg	aggtcaggag	tttgagacca	3420
gcctggccaa	catggtgaaa	ccctgtctct	acc			3453

<211> 3535

<212> DNA

<213> Homo sapiens

60	tcagctcaat	ctttcctccc	tcatagcttt	cttgtcccac	agaagcaggg	tatgggagcc
120	gcaggggcaa	aggagaccaa	tggggaactc	cttcctcaag	agttcctacc	gtacttctcc
180	accaagtccc	ttgtgttttt	gcttggttct	ccccaggtga	gatggtcagg	ctgtacttcg
240	ccatctaaga	ggactctgcc	ctcaccccca	ggctagaggt	gtccaagaat	catgagtctg
300	ctatcttgca	gtaatggact	tccctggag	cctcattgct	ggctctcctg	ctattcctgt
360	cgaagaggcc	cttttgccca	ctctcccata	ctcagctcct	gaggtgttat	ctggactttc
420	tgttcctgag	agggtgtgaa	tctgcccttc	ccttgatgac	ccctctgtac	ataccctgtg
480	aggactatac	gacctgtctg	ggcgtatgag	aggatactgt	ttgtcattcc	caatcggagc
540	acatgatgct	ctgcactgga	tcagagagcc	tcgcactcag	tggaaaggtc	tcagaagaaa
600	cacttagagt	gtttcctagt	taatgattct	cttctttggg	cgtagcatgg	ggaaaatgac
660	actacgaaaa	taaactttt	cttttctttt	ttttatattc	ccattctctt	ctgcatgctg
720	tgcctgggtc	tgtatggagc	agagtcagtg	agagtgaata	gagataagct	tttccaaacg
780	tcttcacctc	atcagcattc	ttgaacaatg	tgcattcage	ggaccttctc	ctcttgaata
840	aatagagagc	agttgtaagg	gagtttttt	ggtttaggta	gacataaggt	attattgatt
900	gagaacccag	ttaaaggaat	agtcataggc	aggagctgtg	attgtaagta	cactcaagtt
960	gcagcctggg	ttgactcagg	aagtagggcc	agactgtggg	tegetagace	aagaaactga
1020	atgtaatgca	tctggtgtga	atgcctgtgt	atgcatgttg	gcaaaagcaa	agcctgaaca
1080	cattttcttc	tcctgctgcc	cttgtactcc	tttctctctg	gtgtatctgt	gctgctcgca
1140	ttcccttata	gtacttcctg	titcctaatt	ttcgtagctc	ttttgtcctt	actgtgaatc
1200	ttcagattcc	catgtatcac	ggcttctcac	ctaatggctt	gctgtgtctc	tttgttgctt
1260	cagattccca	tccttagaga	gttccgaatt	cctacttcca	gtaattttgc	attaccactc
1320	ccatggcaga	ttggcccagg	gggcagagct	catctcggtt	cattactcat	ttgctatagc
1380	ctggtccaat	gtacctggta	cttgggtctg	ataagctgcc	taacctacag	ggctgaagac
1440	tggttgccta	ccacagagca	ggatcacttg	atgatgitca	ccatcatggg	cactigcagt
1500	cttggcagtg	ggaagggctt	gcagtttcct	ctggggcagt	tetgeaaaga	agtgttgctc
1560	gggcttagga	tggaaagggt	tagccaactc	tatagggcag	gattgatgca	caggtgctat
1620	ctgtagcccc	gacatgatgg	taggctaact	aaactttcca	cagacatett	atctttcacc
1680	atgattactc	ttccatggcc	tttggtgttt	gaacccacat	tgattgaaag	aattcagatg
1740	tgatcatttc	aatatttacc	ccaaaaggca	attcagaatt	cagaittigt	tgccttctcc
1800	gatgtccctg	tcaaatgatt	aatttatatt	aaggcatttt	tagataatca	ctttatattc
1860	gggccagtgg	agtgctctga	ctaccgggcc	caactggctc	tgggagtgca	ttatcttttc
1920	tttctcaact	ctgtccagct	ttcaatactg	tgetgtetee	tttagaacte	cateeceact
1980	tttgtacgct	atctctcctt	gicatacaac	ttactgtcct	ccaaatcttg	gatagettet
2040	tccttgtctc	tcccatcttt	ctcaactcat	gtagaaattt	tacactcata	ttectaatca
2100	ttttctgaga	tcccaacatc	gccttccctc	gtatttgcca	ttgacttgag	tttcttatat

accatggaac	catataatgt	catggttgga	agggaactca	aagtttacca	ttttgctaaa	2160
gagacatagt	gtgagagtga	cttatataag	atcacagcat	gctttgcaat	ccagtctctt	2220
gctttccagt	gacccttctg	acttggctgt	gctggctcaa	tatcttttt	ccaccetect	2280
ccactgaccg	tcttcttctt	ccttgtttt	ctatccctaa	gatacaattg	agcattttgc	2340
tcttgttctc	aatgtgtcat	ctcatttctt	aaccataatg	tgacttttt	tcaggtacat	2400
tagctcttta	tgattctgtt	tetteaacag	gaaaagaaat	gagtaaaagt	tgttgtctta	2460
tttttttgg	cagcaggtag	gaacatgatg	gagagttcag	agctgactcc	gaagcaggaa	2520
atttttaaag	gatcagagtc	atctaatagc	acatcagggg	gactctttgg	ggtggttcct	2580
gggggaacag	agactggaga	tgtttgtgaa	gataccttca	aagagttaga	aggacaaccc	2640
tcaaatgaag	aagggagcag	actagaaagt	gatttcttgg	aaataataga	tgaggataag	2700
aaaaaatcca	caaaagacag	atatgaggaa	tataaggaag	ttgaggaaca	tccacctctg	2760
tcttccagtc	ctgttgaaca	tgaaggagtt	ttaaagggac	agaaatccta	tcgatgtgat	2820
gaatgtggca	aagctttta	ttggagttcg	cacctcattg	gtcatcggag	aatccacact	2880
ggagagaaac	cctatgagtg	taatgagigt	gggaagacct	tcaggcaaac	ctcccagctc	2940
attgttcatc	tcagaaccca	cacaggggaa	aagccctatg	aatgcagtga	gtgtggaaag	3000
gcctataggc	acagetecca	tctcattcaa	caccagagac	tccataatgg	ggagaaaccc	3060
tataaatgta	atgaatgtgc	aaaagctttt	aatcagagct	ccaaactctt	cgaccaccag	3120
agaacccata	ctggggagaa	accttatgaa	tgtaaggagt	gtggggcggc	ctttagtcgg	3180
agtaaaaatç	ttgttcgaca	tcagtttctg	cacactggta	agaaacctta	taagtgtaat	3240
gaatgtggga	gagcattctg	ttccaataga	aatctcattg	accatcagag	aacccacact	3300
ggggagaagc	cttataaatg	taatgaatgt	ggcaaagcct	tcagtcggag	taaatgtctt	3360
attcgacatc	agageeteca	cactggggaa	aagccataca	aatgtagtga	atgtgggaaa	3420
gccttcaatc	agatetetea	actigitgaa	catgagcgaa	ttcatactgg	agaaaaacca	3480
tttaagtgta	gtgagtgtgg	taaggcattc	ggtctgagta	aatgtcttat	tcggc	3535

<211> 4208

<212> DNA

<213> Homo sapiens

ć	acaccagcgt	ctcctcaccc	gcaggctgag	acttactgac	acttttcacc	ctcttggccc	60
(cacatgtgtg	gcccactgga	ggggagcaat	ggctggcaga	aaaccaccag	gtctggaggt	120
(caggggctgg	gttctggcct	ggccctgcgt	ctggccaaca	cgtgatctgg	aactgttttg	180
1	tgggcctcgg	ttttccttcc	agaagcagag	aatcattete	catttctgag	tggggcaagg	240

```
gggctctgac ctctcccctg gggctgcctt ttccctttat ttccaagcag tatcacaagg
                                                                     300
                                                                     360
geceaecaaa gtggeatetg gaagtatgtg ttteeaacaa ateeagtett tttttttet
                                                                     420
tttcccctta ttggcttacc atgatatggt agaaattlat tcaggcctgc ctttaaaatt
                                                                     480
ccccttgatt gagtctttgg tggatcttac aaagactttg gtgagttgca gagtctctca
gcatatcacc tggatggagg tagaggggca gagggccacg atgggcagga ttgggcctct
                                                                     540
                                                                     600
gggagaacag aagggccaag gcaaagcaga aaggagcagg gaggtcctcg gcaggggagc
                                                                     660
catctggaag atgicaactt gcaggccctg aggtctgccc tgggctgagt tcctgggtct
                                                                     720
cctactggag agatcagcca aaatacacct ctgttccccc aagcactctt tgctttctgc
                                                                     780
cttctctctg gtcttggaat tttcagttca attgggatct tacaggaaaa caattccagg
ccaattcctg acttcccacc ctagaaggga atgttggagc ccttggcctg tggttccctc
                                                                     840
                                                                     900
teceteette tigggagetg gieeeggggg gaaatggigg gaagiteeat attiggatgg
cccagagtgc tggaatgaga tggagatete cagageagga caagagggte ettatttagt
                                                                     960
                                                                    1020
ctgaagtete caaccigggg citagggtga gagaggaget ggeagaggga tggaceettt
                                                                    1080
aaaggcaaga ggatggcagt tgcaaggaaa ggagctgaga gcttctccga gggtggtaaa
                                                                    1140
aggegtgget tetecagtgg tatecetegg tiletatgit caggacatgi tigaggatge
                                                                    1200
gtccaactga gatgctgaac tccaatattt aaggaalatg atcaaaccga atlaggatta
                                                                    1260
caggacaagt aaagaaaaag aatgcagttg aatctcttga gaggagtatg tgaaagttag
                                                                    1320
aagcaatgat ctcttgaagc tggggcctgt ccctcccact tccttgtctc ccgatggtac
                                                                    1380
taactttata taatactaat aagagtagcc aacatttatt atgcatttgt tgtatgcagg
                                                                    1440
gtattatgcc aagtactica catgicagat cicatiigat teleaceace agettiiggg
                                                                    1500
gcagacatta ttatgttcat ttacaaatga ggacacagag gctggaaagg caaaaaaaaac
                                                                    1560
ttgcaaatgg ctggaaaggg agagagctgg gatttgattc tgggcagtct gactcttcat
                                                                    1620
gaccccagge tetgaactge titaactita tiittattii taaaagiitt aiitattat
ttattittaa ataaatagag acagggtiat gccatclicc ccaggctgat ctcaaactcc
                                                                    1680
tgggctcaag taatcctcca gccttggctt cccactgctg gaactacagg catgagccac
                                                                    1740
                                                                    1800
tgeacceage ceaaactgit ttattetgga tagetggaca tageteecet etgagattaa
                                                                    1860
caggacgaaa tatgaggeec acageettte cageteagae cattgeaate ectagtteet
                                                                    1920
teegtteeae tetetettae eecaggagte aggaagttet tetgtgtetg acceteaett
                                                                    1980
cttcctgtgg taaagtgatt tctgagatta gaatcttcaa alcctcttgt gattagcatg
                                                                    2040
ageigeatee catticaaag eecaceatgi igggeactia ataaatgita aacattiett
                                                                    2100
atggggtate acateacetg catetecece cattggetee etgecaacat geacagatag
                                                                    2160
tetettetgg attecactgt tgaaggeagg aaaactgeee ttettttett ttteeteete
                                                                    2220
ecceatiggg acatatggic ictgitgagg geagagigag gaaccicaac aggegicetg
                                                                    2280
tggcttgggg tcagcctagg ctgttgcctc tgaaccctgt gttgcactgc agcctgagga
                                                                    2340
agtttagggg tgcagaggac caaaggagac acatatggat tcctggaaga ggaattggtg
                                                                    2400
ttggccagca titclagagi cggalaigca galeccalee cicigicele ccaeccalig
```

gccatcccac	ctgcctccat	ttgtagacag	ttcctgaaag	atcaggatat	tatgtatgca	2460
aagaggccag	agagcttcag	ctgtaagaag	cttgcaatga	gcaagaaggt	caccccagct	2520
tcaactcaat	gcgtccgctt	cccttcccc	tccacagtgg	tctgcaagaa	gaatctggac	2580
agtaccactg	tggccgtgca	tggtgaggag	atttactgca	agtcctgcta	cggcaagaag	2640
tatgggccca	aaggctatgg	ctacgggcag	ggcgcaggca	ccctcagcac	tgacaagggg	2700
gagtcgctgg	gtatcaagca	cgaggaagcc	cctggccaca	ggcccaccac	caaccccaat	2760
gcatccaaat	ttgcccagaa	gattggtggc	tccgagcgct	gcccccgatg	cagccaggca	2820
gtctatgctg	cggagaaggt	gattggtgct	gggaagtcct	ggcataaggc	ctgctttcga	2880
tgtgccaagt	gtggcaaagg	ccttgagtca	accaccctgg	cagacaagga	tggcgagatt	2940
tactgcaaag	gatgttatgc	taaaaacttc	gggcccaagg	gctttggttt	tgggcaagga	3000
gctggggcct	tggtccactc	tgagtgaggc	caccatcacc	caccacaccc	tgcccactcc	3060
tgcgcttttc	atcgccattc	cattcccagc	agctttggag	acctccagga	ttatttctct	3120
gtcagccctg	ccacatatca	ctaatgactt	gaacttgggc	atctggctcc	ctttggtttg	3180
ggggtctgcc	tgaggtccca	ccccactaaa	gggctcccca	ggcctgggat	ctgacaccat	3240
caccagtagg	agacctcagt	gttttgggtc	taggtgagag	caggcccctc	tececacace	3300
tegececaca	gagctctgtt	cttagcctcc	tgtgctgcgt	gtccatcatc	agctgaccaa	3360
gacacctgag	gacacatctt	ggcacccaga	ggagcagcag	caacaggctg	gagggagagg	3420
gaagcaagac	caagatgagg	aggggggaag	gctgggtttt	ttggatctca	gagattctcc	3480
tctgtgggaa	agaggttgag	cttcctggtg	tccctcagag	taagcctgag	gagtcccagc	3540
ttagggagtc	actattggag	gcagagaggc	atgcaggcag	ggtcctagga	gcccctgctt	3600
ctccaggcct	cttgcctttg	agtctttgtg	gaatggatag	cctcccacta	ggactgggag	3660
gagaataacc	caggictiaa	ggaccccaaa	gtcaggatgt	tgtttgatct	tctcaaacat	3720
ctagttccct	gcttgatggg	aggatectaa	tgaaatacct	gaaacatata	ttggcattta	3780
tcaatggctc	aaatcttcat	ttatctctgg	ccttaaccct	ggctcctgag	gctgcggcca	3840
gcagagccca	ggccagggct	ctgttcttgc	cacacctgct	tgatcctcag	atgtggaggg	3900
aggtaggcac	tgcctcagtc	ttcatccaaa	cacctttccc	tttgccctga	gacctcagaa	3960
tcttcccttt	aacccaagac	cctgcctctt	ccactccacc	cttctccagg	gacccttaga	4020
tcacatcact	ccacccctgc	caggeceeag	gttaggaata	gtggtgggag	gaaggggaaa	4080
gggctgggcc	teaccgctcc	cagcaactga	aaggacaaca	ctatctggag	ccacccactg	4140
aaagggctgc	aggcatgggc	tgtacccaag	ctgatttctc	atctggtcaa	taaagctgtt	4200
tagaccag						4208

<210> 4680

<211> 4201

<212> DNA

<213> Homo sapiens

<400>	4680

60	tgtattaata	cttcacgctc	tgtgcagcac	tgatattctt	attactcaga	tatattattc
120	cgccccagaa	gcttggttat	acagcagctg	gcaagtggtc	ctgcattaca	tcttgttttt
180	taaggtcatt	agctagagga	ttagagatgc	ttatcactat	cttcatctcc	aaaattcaca
240	tttccaaaaa	ctctaaatga	tctttaaaaa	cagacgagac	aagttcaact	aagcctcaaa
300	tgctgtgtct	ttcctactta	tccctacgca	gattcgtccc	atattaattg	ttatttggag
360	tttgactcct	gtaaaaggtc	gatttacgca	gggcaatcct	caatattacg	aatctttttg
420	ggttaccaga	aacagtçtca	aaatgcattc	acttataaaa	ctaaattacg	gaggcagatt
480	accaacagga	ctgaacattc	atttttccca	tgaagtacta	atttaccttt	gtaaatccac
540	ctctttacgt	tcccacacag	ttgtgttttc	tttaattgaa	agggacataa	ctcattattc
600	cgtctccttc	ggctcgttct	taattggtca	atttctacct	tctagatcaa	acactaacta
660	caagttcaac	tacccgatta	ttgttcccct	caaaaaaatcg	aacagagcct	gtctttcagg
720	gtataattga	cgtttcccag	gcatttggca	ctiggcaagi	acttgcattg	aggctttgct
780	ggattttacg	ctcacgtctt	gttccttaaa	aattatttca	cctcatgtaa	taatcactat
840	atgcttcctc	gttttcactg	agctgtcact	cattacctga	agaaacaccc	taacattact
900	gggctatttc	cttaacacag	agaacgtgtt	caggacccag	gcagcatata	taatggccga
960	ctgagccagt	gaagattttc	ggctgtcctt	ctgctgttat	gctgaactgc	agcacaacga
1020	aaactgcctt	cgcaacattg	acatgttgct	cgtatgtcgt	tctgactcag	taacattttc
1080	agtctgtgct	caaaagtttc	ccttctttt	ataacctatt	ctgcctgatg	aatcaaattt
1140	ccctccccgg	gcccacacac	tcacattcgg	tctacagtac	tcttctcctt	cagagcaagg
1200	ttacagatgc	actcccgttt	tacattagtt	ccagagctga	gcagcaaatg	acccctctcg
1260	agttccccct	ctccgaaaca	tgctgcggga	ctcacgtcaa	catgctttaa	agaaaatttt
1320	tgttaatttt	acttgtcaag	tcactgtcct	ctactgtgcg	caagctaaaa	tacatggaag
1380	ttiggcaaai	cagaatgctc	aggactttca	ttaaccctag	tcttcaggag	accaccactg
1440	ctattgatac	atacatgtaa	gctttctttt	cttttggcaa	cattatecte	ggatgtgact
1500	atggtaaacg	agtacagctc	aacgggcgaa	ccacttgtga	tttatttggg	cttttctcat
1560	ctgataacgg	aaactaaaaa	ctgttctgag	cggttatggg	tcctgtttct	acatgtgcct
1620	ttactcacac	acatgggcaa	gtttcttcag	cttttaaaaa	actagtgctg	ccccgcctac
1680	ataaaacact	gaacgagcca	ggccttggtt	ctggagaaca	ccctataact	aacgggcatt
1740	ctcctcatgc	gatgctatta	atcaaaggga	aggatactaa	cttcgtaaac	caaagatcaa
1800	atcaaccttt	ttaacaagaa	tttcttaaac	tcacactaaa	ctggctcttt	ccaattaagt
1860	gaatgcgggt	ccacaaaaag	taaatttgat	tcactggaaa	gaacaacatt	cacageegea
1920	cttggggtag	acggtcataa	ggaattaggc	ctaacacatg	gacgtaaaaa	atggtggaag

```
gggttttgtt tgtgtttccc caggaaaaga actacagcct gtgtgggtcc cctcccgaca
                                                                    1980
                                                                    2040
attaaaattg caccataact ccaaagatga aacgctccca gaaggaaaag gcaaagaccc
                                                                    2100
ateggaaaca aaagagcaac tttcaccgcc tgatacatac acttcatgac atccacattg
                                                                    2160
agacaaaatc tcgccgtgtg acctacaaca cctctcgttc aaccccacct acttggggtc
gaatacagat tttatctcat caaacagaaa aattcttaac agaaaaagga atcccaaaaa
                                                                    2220
                                                                    2280
tgactggtaa tataattctg gctgccttta tggtagtcag tgcagcggta agtataccac
cggtcgggc aactcaaaat tatacttatt gggcatatgt tccttttccc cttctaattc
                                                                    2340
                                                                    2400
ggtctgtctc ctggagggac tctccagtag aagtttacac taataatagt gcattcatgc
                                                                    2460
cgatccctaa tgatgatcgg tttccagctc aaccggaaga agaaggtatg cactttaatc
                                                                    2520
tgtcaattgg ctataaatat ccaccattat gtattgggaa gtcgcctggt tgtttagctt
attettaaca gaattggatg tggactgtac cgtcctttac aaatgattet tateaagtat
                                                                    2580
ataatgtgtt cagtactaac tcttttcaac ttctcactgt caaacgtacc ccacatgagg
                                                                    2640
catggagagt tecteteact accaaaacta ataaaacaaa aggactgeeg gactgteeaa
                                                                    2700
                                                                    2760
agaaacctac aaatgggcct tttatagtga cttcaatttt atgggataat tgtaatgctc
ccaaggctgt tgtactccaa actctagcca tgggtattgt tattgattgg gctccaaaag
                                                                    2820
                                                                    2880
gacattattg gcaagattgc tccagcaaaa ataccttatg ctcggagttt atttattcct
                                                                    2940
tagattatat agagcatggg tggcagtctt acacgatgag acaacgggtg tctccttacc
                                                                    3000
catttaaatg gatggacaca ggtattgctc ctcctagacc aaaaattatt catccctttt
ttaccccaga acatcctgaa ctatggaaat tagctgcagc tttgtcggga ataaagatat
                                                                    3060
                                                                    3120
ggaacactac ctatcagete ettegtaeta aaaccaaaac acceacatte aacatcaece
ttatttctga atgggtgata cccattagga gctgtgtcaa acccccttac atgctgttgg
                                                                    3180
                                                                    3240
ttggaaatat aattatgatg cetgatgeac aaactataga atgteataac tgtaagetgt
                                                                    3300
teactigeat igaigeaaci titaateeea etacaagiai teletiggia agggelaggg
agggggtatg gataccagtt tetetacate gtecatggga gtettecece tetatteaca
                                                                    3360
tagicaatga agitettaaa gacaleetea aaagaacaaa gagattiatt iitaeletta
                                                                    3420
                                                                    3480
ttgcagtcct tgcaggacta cttgcggtta ctgcaacagc agcaactgct ggagttgcca
tecgeagtte tgtteaaact geteactaig tigaageaig eeagaaaaat teeteeagae
                                                                    3540
                                                                    3600
tttggaatte teaggegeaa attgateaaa aattagetaa teagattaat gateteegee
                                                                    3660
aaagtgtaac ctggctggga gatagagtta tgaacttgca acaccgtatg caattacagt
                                                                    3720
gtgattggaa tactictgat tattgcataa cgccttatgc ttataatcaa gatcaacata
                                                                    3780
gctgggaaaa tgtctcaaga catttaaaag cctgggatga taacttaacc ttggatattt
                                                                    3840
cacaacttaa agagcaaate tttgaggett cacaagteca tttatecaca gtteetgget
cacacattti tgaaggcata actaaacaat tacctgattt taatcccttc aaatggctca
                                                                    3900
                                                                    3960
aaccegicag aggateatig tigitacigg cattatiaat atiggiatge tiatgilgic
teettttagt etgeagatge etetaaggag teegaaacea agteegaagt caacaacaag
                                                                    4020
                                                                    4080
caatgatggc gatggcgatt ctagttaata aaaagggggg agatgtgggc ggcaagccac
```

ccaggcaccg	aggcaagaga	cagaggacac	gagctgtacc	agtataataa	aatataaaac	4140
aagaatagtt	ataccagata	tagatcttag	atatgattat	atacgaatat	cattaatcat	4200
g						4201

<211> 3847

<212> DNA

<213> Homo sapiens

			*			
60	gtctgtgtca	cgtcagctca	acgagggccc	cccacagage	ccccacccca	aggcccctga
120	ttcctggaca	ccacctgggc	cctcctcggg	ctgtctgcca	cctccgtgtg	gccccgatgg
180	ttggccctcg	cgccccggtg	gctcccacac	atgctggctc	ggtgtaccac	cgctgtcccg
240	gtccgcatct	ggaccgtacc	ccgtgtccca	cagctggcca	gaggcgggga	ccatggagca
300	ggaggccgct	ccttcagcct	gccgtgcgct	cagcagtggg	caccctgcaa	ggggcctggc
360	cgccacccct	ccggcctgac	ggagctgtca	gtgccaccga	tggaacacac	gaggtcctgg
420	cagctgtgcg	tggcccagta	cagggctccc	ctcctgctcc	tgctcttcag	gacggccgcc
480	tgccccgcg	tatgcccgga	gcggacatgg	ccgagtggca	ggcatgtcct	gacccccagt
540	gggaccctcc	tggcctttgt	ggccgcctgc	cagcagggat	ccctggcagt	agccccagcg
600	agttgacatc	agctgctgcg	tcccttgatg	gggctcggcc	tgacagtcat	aggtgcacag
660	ctttggccct	tggctgtgtg	gactcagcca	cagccgcctg	acctggccag	ggcactctgg
720	gctggatgct	gagtcgtggt	tcgtccaaca	ggtgtccacc	gccaactgct	gcagctctgg
780	ccctccttg	ctgagccctg	ggtgtccacc	ggagctgccc	gcatcatccg	gtgtcgggcc
840	caaggtgtgg	gccggaccat	attgccgccg	cttcctgctg	aggacgcccg	acgctcagtg
900	ggaacccgtg	teggeeacte	caggtgtaca	cccaggcccc	cacaggccag	gactacgcca
960	cgccgtcttc	gcgcagggga	caggtcctca	tgaccagcag	ccttctctcc	caggctgtgg
1020	cccccagcc	tccccggggc	gaccaaagct	tactgagagc	tcctggccac	ctctgggatg
1080	cagegagete	cgtccagggc	gaggacgcag	aggaccgctg	gcccgggcgc	tgcaagacag
1140	cgtctgtgcc	cacggctggg	gcatctccac	gccatgtcag	aggtccccaa	ccccggcagc
1200	cccacgcacc	attcgggggc	gacaccagga	tggcgccagg	aaggtggcga	aggcctcccg
1260	ccccactct	tcagctgcag	cttgccaggg	ggccttcacg	cttcctgcaa	acctacctgg
1320	ggctgtcgtc	tgcgtctgaa	ggtgggtggc	tcccgccagc	cttgcccgcc	gccaagggca
1380	tgggggccaa	cggacacagg	gtctggaggc	ggccaacatg	ggaatgggcg	ggttacagcg
1440	ctccacccca	gtccagccgt	gccgcgcgtg	ccaagatgcg	ccacccccag	gagcctaccc
1500	ttctcccggt	tggggctctg	gccatcagtg	atgagcgcca	ctggtggcaa	aggccaggac

agcgcctcct	ggcgctccag	ccgaggcttc	tttgcctaca	cgtgcggccg	cctggtggtg	1560
gtggaggacc	tgcactctgg	cgcccagcag	cactggtccg	gccactctgc	ggagatctcc	1620
acgctggccc	tcagccacag	tgcccaggtc	ctggcctctg	cctcgggccg	aagcagcacg	1680
accgcccatt	gtcagatccg	cgtctgggac	gtgtctggcg	gcctctgcca	gcatctcatt	1740
ttcccccata	gcaccaccgt	gctggccctg	gccttctcac	cagatgacag	gcttcttgtc	1800
acactggggg	accacgatgg	ccgcaccctc	gccctgtggg	gcacggccac	ctatgacctc	1860
gtgtcctcca	cccgcctccc	ggagccggtg	catggtgtgg	ccttcaaccc	ctgggacgcc	1920-
ggtgagctca	cctgtgtggg	ccagggcact	gtcaccttct	ggctccttca	gcagcgtggg	1980
gcagacatca	gccttcaggt	gcgtcgagag	ccggtcccag	aggcagtggg	ggctggagag	2040
ctgacctcgc	tctgctacgg	ggcacctccc	ctgctctatt	gtggcaccag	ctctggccag	2100
gtctgtgtct	gggacacgcg	tgccggccgc	tgcttcttgt	cctgggaggc	ggatgacggt	2160
ggcattgggc	tgttgctgtt	ctcgggttct	cgattggtca	gcggcagcag	cacggggcgg	2220
ctgcgcctgt	gggccgtggg	ggctgtgtcg	gagctgaggt	gcaagggctc	aggcgccagg	2280
tccagttctg	tgttcatgga	acacgagctg	gtgctggacg	gggctgtggt	gagtgccagc	2340
ttcgatgaca	gcgtggacat	gggcgtcgtg	ggcaccacgg	cgggcacgct	gtggtttgtc	2400
agctgggccg	agggcaccag	cacacgtctc	atcagtggcc	acaggagcaa	ggtgagggac	2460
ttccagcctg	ggcggaggcg	gggcagccga	acctggtgcc	ctccctgcct	gccggctcca	2520
tctccaccag	cccagatgat	tccaagtcct	gccgtcactg	gctcgcagcg	gccgccttgg	2580
ggttcccagc	ggggaagtct	tgggtgtgca	cgtcccctca	aagccgtccc	ggttgtgtct	2640
gcacaagcga	gccgcctggc	aggccttgca	ggtgaacgag	gtggtcttca	gccccgggga	2700
glcccactgc	gccacatgca	gtgaggatgg	gagtgtgcgg	gtgtgggcct	tggccagcat	2760
ggagcttgtg	atccagttcc	aggtgctgaa	ccagagctgc	ctctgcctgg	catggagccc	2820
cccgtgctgt	ggccaccctg	agcagcagcg	gctagcggct	ggctacggtg	acggctccct	2880
gcgcatcttc	agcgtctccc	gcacggccat	ggagctcaag	atgcaccccc	acccggtggc	2940
gctgaccact	gttgccttct	ccaccgatgg	tcagactgtc	ctctctggag	acaaggatgg	3000
gctcgtggct	gtgagccacc	cctgcacagg	gacaaccttc	cgtgtgctga	gtgaccacca	3060
gggcgcccca	atctctacca	tctgtgtcac	gtgcaaagag	tgtgaagact	taggggtgga	3120
gggcacagac	ctatggctgg	ctgccagtgg	ggaccagcgg	gtcagcgtct	gggcctccga	3180
ctggctgcgg	aaccactgtg	agcttgtgga	ctggttgagt	ttcccaatgc	ctgccaccac	3240
ggagactcag	ggccacctgc	caccctccct	cgctgccttc	tgcccttggg	atggggcgct	3300
cctgatgtac	gtgggccccg	gtgtttacaa	ggaggtgatc	atctacaacc	tctgccagaa	3360
gcaggtggtg	gagaagatac	cactgccctt	ttttgccatg	tccctgagcc	tgtcccccgg	3420
gacccacctc	ctggctgttg	gctttgctga	gtgcatgctg	aggctggtag	actgtgccat	3480
ggggactgcc	caagactttg	ccggccacga	caacgcagtg	cacctgtgca	ggtttacacc	3540
ttccgccagg	ctgctcttca	cggccgcccg	caacgagatc	cttgtgtggg	aggtccccgg	3600
cctctgagat	gcagcaggga	ctgtggtggt	gggcatcacg	cctggtcatg	ccaggcacct	3660

ggacacaggc	ttggcagagg	${\tt cgccaggttg}$	tcaatggcct	catgctggga	caggccagga	3720
ttcacgtaaa	tcgcctggag	caagctgttg	taaatttggc	gccctgtgaa	tactttcata	3780
cctgttgccc	ttttgcctaa	gaaatcttta	atgtttctat	cttgtaataa	acatgggcat	3840
ttatttc						3847

<211> 3913

<212> DNA

<213> Homo sapiens

<400> 4682

acttattcct gcaaatgtga attaatcctt ttcttacttt gagcatgacc atataacttg 60 120 ctttggtcca taggacatta gcaagagagt gcaaaccaag acttcataag tacttgtgca ttggagcttg gctatctaga aagttccctt tcgggagcca gttactatgt tgaaaagatg 180 240 cttgggctga atcatgagag gcaacatgga gagaggcatg gggaattagg agacccacca 300 gctgaaccct gccagaattt ctgagaaaca gaattctgag aaataataaa ttgttgtttg ggggtggttt gttatatggc aatggaagac tgaaacaagc tataacagcc ttcacaacaa 360 420 actitettet tiettiaaac aaacaaacta tgeattaact tagtietgat tatteetitt 480 taaactcctt ccctagttit cctcttatig tctcaacttc gtcttttctc atggaggtgt 540 ggatggctgg tgctataagc attgtgagtt gggagatact ggcaagccct gcttaccaag 600 aatettttee ttateageet gleetggtae tigaaacite aceattatat aaataaggaa 660 aattttetee taaactaaag aatttteate teeaagaata aatatgeaac taleeetgga 720 cacatacact attctttgtc attcaccatt gagccaggat taatttgtaa gtttgctttc ctgggaaaca gctaactgag aaggaacaga aagacttgcc cacacattga cttctgattg 780 840 gtaggetett eeeetttte tetateeaaa taaeteetge tetttettea getettttt tttttgggag acggagtete getetgtege ecaggetgga gtgeagtgge aglggegega 900 960 teteggetea etgeaacete egeegeeegg gttegagtga tteteetgee teageeteee 1020 gagtggggtt accgcaggca tgaaccacca tacctgccta gtttttgtgt ttttagtaga 1080 gacggggttt tgccatgttg cecaggctgg ttttgaactc atggcctcaa gtgatccatc tgccttggcc tcccaacgtg ttggcatgag ccaccatgcc tggcctgttc ctlcagcttt 1140 1200 taactccetg aaccetgtga ttaggttaat tttcccatta taggetetga tagtaccaaa anacteetet tigageaett ategeagigg caatteaace titittaigi glaattalat 1260 1320 ttatatetee tagaactgta ageteeaaag ggteagagae ettgtetgtt ttgaceaetg tagtattece aattetteee atataegetg acacacagaa tgtactcaat attigttaaa 1380 1440 tgaatgagtg aatgaatgaa ctgtgattct gaacaattaa attctctctt tetttettta

gaaattcttg	tgcctggact	tggtttttt	attactattg	ggggcaagaa	agcagtcact	1500
tggcattcta	aaagctctat	gagatgtttc	aagaggagga	aattaaagct	aagtgtgcta	1560
tgcttgaaat	gaagacttat	tagaccaatg	taggctccct	tacaatgcct	gacccattic	1620
agttgtagcc	catgaacaaa	atcatttcca	gaagtgagaa	aactaataag	agctaacatg	1680
ccttgtacta	gcaggcattg	taaaaacatc	actggagcgc	cgtcaatatt	attcctgttt	1740
ttccacacca	taataattta	tagcttccct	ttcttttata	attaaataat	ttcagacgtg	1800
ttaaggaaga	accataatgg	tttaatacac	tttgcagttc	tctgatttat	cttcactgaa	1860
gcctgtaata	attaaacata	aaaattatat	acagtgggtt	tattaatata	agaaaaaagg	1920
aagtgatgga	gctttagatt	ttcttttaat	attgctattc	cacctttgaa	tgggttgagc	1980
aagaatagca	atcatgtttt	aaccaatgtt	tatcttatga	atagtatttt	tgtcattttc	2040
tgaggggtgt	gtgtcaaaag	ctgccacttt	tatttcccta	gggataattg	ctgcttgttc	2100
ttcaatttag	aatatctggg	gtagtaattg	atctggagaa	gcacttctta	aagtgcaatg	2160
ctaattgcag	gaaagagaaa	aatgggttct	gtggtcacac	aggtttggga	aatgctgcct	2220
taaataaaat	taaatagtgc	tttttttttt	ttgtcacagg	attgataata	atatgctctg	2280
taaatctcca	aaaggaaggt	tataatatac	agcatttccc	attgtcatct	gatcattgaa	2340
ttttaaattt	aattaaacac	ttttttggtt	tggtatgttc	tttgacattt	aaatgtgttc	2400
tctcagttta	aattttggga	gacttaaccc	tcaaaatgtt	tgtgagatat	aaataatcct	2460
cttcttatgc	atgtagagag	gtgtatacag	gcataacttg	gagatattgc	aggttcagtt	2520
ctagaccacc	acaataaagt	gaatatcaca	ataaaatgag	tcacacaaat	gttttggttt	2580
cccagtgcat	ataaaagtta	tgttggtggg	aggagccaag	atggccaaat	aggaacagct	2640
cctgtctaca	gctcccagcg	tgagcgacgc	agaagacggg	tgatttctgc	atttccatct	2700
gaggtaccgg	gttcatctca	cttgggagtg	ccagacagtg	ggcgcaggtc	agtgggtgcg	2760
cgcaccgtgc	gtgagccgaa	gcagggcgag	gcattgcctc	actigggaag	cgcaaggggt	2820
cagggagttc	ccittccaag	tcaaataaag	gggtgacaga	cgcacttgga	aaatcaggtc	2880
actcccaccc	gaatactgcc	cttttccgac	tggcttaaaa	aatggcgcac	cacgagatta	2940
tatcccgcac	ctggcttgga	gggtcctacg	cccatggagt	ctcactgatt	gctagcacag	3000
cagtctgaga	tcaaactgca	aggcggcagc	gaggctgggg	gaggggcgcc	caccattgcc	3060
caggcttgct	taggtaaaca	aagcagccgg	gaagctccaa	ctgggtggag	cccaccacag	3120
ctcaaggagg	cctgcctgcc	tctgtaggct	ccacctctgg	gggcagggca	cagacaaaca	3180
aaaagacagc	agtaacctct	gcagacttaa	atgtccctgt	ctgacagett	tgaagagagc	3240
agtggttctc	ccagcacgca	gctggagatc	tgagaacggg	cagactgcct	cctcaagtgg	3300
gtccctgacc	cctgaccccc	gagcagccta	actgggaggc	acccccagc	aggggcacac	3360
tgacacctca	cacggcaggg	tactccaaca	gacctgcagc	tgagggtcct	ctctgttaga	3420
aggaaaacta	acaaatggaa	aggacatcca	caccaaaaaac	ccatctgtac	atcaccatca	3480
tcaaagacca	aaagtagata	aaaccacaaa	gatgggaaaa	aaacagagca	gaaaaactgg	3540
aaactctaaa	aagcagagtg	$\tt cctctcctcc$	tccaaaggaa	tgcagttcct	caccagcaac	3600

agaacaaagc	tggatggaga	atgactttga	cgagctgaga	gaagcaggct	tcagacgatc	3660
aaattactct	gagctacggg	aggacattca	aaccaaaggc	aaaaaagttg	aaaactttga	3720
aaaaaattta	gaagaatgta	taactagaat	aaccaataca	gagaagtgct	taaaggagct	3780
gatggagctg	aaaaccaagg	ctcgagaact	acgtgaagaa	tgcagaagcc	tcaggagccg	3840
atgtgatcaa	ctggaagaaa	gggtatcagc	aatggaagat	gaaatgaatg	aaatgaagca	3900
agaagggaag	gtt					3913

<211> 4376

<212> DNA

<213> Homo sapiens

<400> 4683

60 tgaccttctc tcctccctgt gtcttttttt ttcctttaga gatgggggtc ttgctctgtc 120 actocagetg gagcacagtg gtgcaatcgg agctcactga etgcggcetg caactoetgg 180 gctcaagtga ttctgccgcc tcagcctctc aacgtgctgc gattagagag tctgtagaag 240 aggogacacc agggcttcca aatgaacaac cgaaaggaag atatggaaat cacgtccac 300 taccggcacc tgctgcggga gctcaacgag cagaggcagc acggcgtcct gtgcgacgtc 360 tgcgtggtcg tggagggcaa ggtcttcaag gcgcacaaga acgtcctgct gggcagcagc 420 cgctacttca agacgctcta ctgccaggtg cagaagacgt cggagcaggc cacggtcacg 480 cacciggaca tegicaegge ceagggette aaggecatea tegaciteat giacteageg 540 cacctggcgc tcaccagcag gaacgtcatc gaggtgatgt cagccgccag cttcctgcag 600 atgacggaca tegtgeaage etgecacgae tteateaagg eggegetgga cateageate 660 aagteggaeg ceteagatga gettgeggag ttegagateg gegeetegte eageageage 720 acggaagete teatetegge egtgatgget gggaggagea teteecegtg getggeaegg 780 cgaacgagtc ctgccgattc ttccggagac tcggccatcg ccagctgtca cgacggaggg 840 agcagctacg ggaaagagga tcaggagccc aaggccgatg gccctgatga tgtttcttca 900 cagectetat ggeetggaga egtgggetae gggeetetge geateaagga agageaggtt 960 teacegtete agtaeggagg gagegagetg cettetgeea aggaeggtge agtaeagaae 1020 tettteteag ageagagtge tggtgatgee tggcagecea egggeegaag gaagaategg 1080 aaaaacaaag agaccgtccg gcacatcaca cagcaggtgg aagatgacag ccgggccagc 1140 tecceggige egiecticet geogaegieg gggiggeegi teageageeg agaeteaaat 1200 geggacetgt cegteacega agecageage teegacagee gaggagagag ggeegagete 1260 tatgcacagg tggaggaggg tctcctggga ggagaagcca gctatctggg ccctccctc 1320 accccagaga aggacgacgc cctgcatcag gccaccgcgg tggccaacct gcgcgcggcg

ctcatgagta	agaacagcct	gctgtcgctg	aaggccgacg	tgctggggga	tgacggctcc	1380
ctgctgttcg	agtacctgcc	cagaggggcc	cactcgctgt	ccctgaatga	gttcacggtg	1440
atcaggaaga	agttcaagtg	tccgtactgc	agcttctcgg	ccatgcacca	gtgcatcctc	1500
aagcgacaca	tgcgctcgca	cacgggagag	cggccctacc	cctgcgagat	ctgcgggaag	1560
aagttcacgc	ggcgcgagca	catgaagcgc	cacacgctgg	tccacagcaa	ggacaagaag	1620
tatgtgtgca	aggtgtgcag	ccgcgtcttc	atgtccgccg	ccagcgtggg	catcaggcat	1680
ggctccaggc	gccacggtgt	gtgcaccgac	tgtgctggcc	gcggcatggc	cgggcccctg	1740
gaccatggcg	gcggaggcgg	cgagggctct	ccagaggcgc	tgttcccagg	cgacgggccc	1800
tatctggagg	accctgagga	cccacgaggg	gaggcggagg	agctgggcga	ggacgacgag	1860
ggcctggccc	ctgaggatgc	gctgttggcg	gacgacaagg	atgaggaaga	ctcgccgcgg	1920
ccgcgcagcc	ccccaggagg	ccctgacaag	gacttcgcct	ggctctccta	ggcccgcccg	1980
ccggcagggt	cggtggctgc	ctcgctccgt	ccacccgtgt	gtgtgtccgg	tgggtctcca	2040
ctgcggggcc	agggccacgc	tcacccctct	cgcggcccct	cctctgcttc	cccctgaacc	2100
caccccccac	ggaaaccagc	cctgcgggct	aagcaggtgc	gaccccagca	agaggggtgc	2160
tctgggacca	gacatgaagt	gagttggggg	agggcacagg	gtgggtttga	gtgaagggag	2220
agcacggtcc	taagtcccca	gcaggtggtg	cgggtgtgtg	agtggcccct	gtgatggcca	2280
gcctggcttg	gacacgtgat	gggcctgtgg	ccgggtccag	tgggcactgg	gcggggtggt	2340
gtgttcggcc	cagaggcccc	tggcctgagc	aggtgcagag	ttttacagac	acccggtcag	2400
cccggctgga	gcccgccctc	ccctccccta	gcagccaggt	cactgcctgt	ggctgcagcc	2460
gtggcccgtg	gtgcagccgt	ggcccatggc	agcctgtaga	atccaggtgc	acagagagec	2520
ctgccaccgt	cttacctctg	ggctttggtg	cttaacacac	aacacagctg	cagaccctgc	2580
tggaggccga	gggctccagg	tgctatctta	ggtggacaca	gccctggggg	ctccttccag	2640
gaggggaccc	tcagccctgt	gcccccacc	acttcaggcc	acaccaggtt	ccctctgcaa	2700
gggcctcggc	tcagtcgtgt	gcacttcctc	ggagagcctt	gggctgccac	gtccaccccg	2760
ggctctgccc	gtcctgttct	gcccatggcc	cagcccggcc	gctcctgctg	acccctcctg	2820
gacgggttgg	agctgggctc	ctgcctttgc	tgctaacact	ggaggcggtg	ttcctaactg	2880
cagtgtgctg	cttacacctc	cccgcgtggg	taaccaaatt	tttaagtagt	cagagacata	2940
tcgaggtagt	tacataaaat	tattttgttt	ggcattattt	ttctcactcg	aagaaactat	3000
atagggttgt	ttttccttta	gcttgtgctc	aagtcctctt	gctgtgtttt	cagaagcact	3060
cacatgttct	ttcttttcct	gagtgaaaag	caaaggtccc	acggtgtgtg	ctgtggtgca	3120
ccgcctggct	ttgggggtcc	cggaggcagg	ctgcctagac	tcacagcctc	gggaccgttg	3180
ccacggcctg	tcttctcgtt	caggcctgcc	tctgacagca	ctcaccatga	ggacattcca	3240
tectteacce	cctcctctgg	cacaggccac	cactgcggtg	ctgtgccttc	agatgggagg	3300
tgggcgcggt	ggccgcctcc	ttccctccag	gacctgcccg	tgtgaagacc	ccccggagtg	3360
ctgagcttca	gggctgcgtg	gaaagagttt	ttactctctt	tttctagcct	gtataccagg	3420
cttttcccca	cattgtcagg	tagagcacca	gcttccctga	ccgctgctgc	tcggggaggg	3480

ctggggctgg	ccgggggtcc	tgtggaggag	tacatggagg	actccaggta	cagcgcagga	3540
gtcacggctt	ttgttttttg	acattggccc	ccggttctac	caatgacagg	gtgccctggc	3600
tggagctgtc	atcacacaca	ccctcagct	cggaggctgt	gggctcctca	aagctggaga	3660
aagaggccaa	gatttttctg	cacacggagt	gtggggatag	gagccgggcc	aagcgctggc	3720
ccctcagcgg	tgagccctgc	ccactcttac	cgagcaaggt	gggtggctct	ggcacgagtc	3780
ccccaggggg	agagcatggc	taccagggag	ctgcagcgga	gccctccagc	cctcacccca	3840
ggccagcccc	accccggcct	ctttgagaat	tctcagaact	ttgtaccttt	cccctgattt	3900
ttaaaccctt	tttctaaaca	gactgacttt	cttacaaaat	gcatttggaa	accagacctt	3960
tgctacccac	caatgtctct	gggttttgta	ccagtccctg	ctctcaggcc	accetgecea	4020
ggacccaggc	ccgcctcccc	ctccacactc	aggatgtcct	gctccatctg	gccggctcac	4080
tccgtgtggc	ctgcctttgc	tgaccgtttt	ggggttcccc	gccggagcta	caggggcatt	4140
ttcttcccta	aaaccaacag	tgtcccactg	acctccccaa	gtgtttgctg	cgtggcagat	4200
ttcctgttct	tgttcgcagt	ttgccgactg	aagagtgtgg	gatttccgag	gcccaggtga	4260
gcacgtccat	ctcaggaggc	gtggagggaa	aagacatgtc	atgaagggtt	ttttttatgt	4320
gactgatttt	tttttaaatc	gatgttcaaa	ctaataaata	tttttttatg	aagagg	4376

<211> 492

<212> PRT

<213> Homo sapiens

<400> 4684

Met Pro Ser Phe Leu Val Pro Ser Leu Val Ser Ser Pro Val Leu Leu l 10 Lys Leu Leu Phe Ser Pro Gly Pro Lys Thr 11e Trp Ser Leu Trp Gln 25 Gln Pro Met Leu Phe Gln Glu Ala Thr Ala Phe Glu Asn Met Thr Lys 35 40 45 Asp Trp Asn Tyr Leu Glu Gly Ser Gln Lys Asp Cys Tyr Arg Asp Thr 55 Met Leu Asp Ser Tyr Glu Asn Thr Val Pro Gln Gly Ser Phe Leu Gln 65 70 75 Leu Ser Met Met Pro Gln Arg Ala Gly Asn Asp Pro Pro Gly Val Ser

85 90 95

Asn	Ala	Ser	Glu	Met	Glu	Met	Glu	He	Ser	Asn	Met	Arg	Glu	Lys	Phe
			100					105					110		
Leu	Met	Ser	Val	Thr	Lys	Leu	Val	Glu	Ser	Lys	Ser	Tyr	Asn	Ser	Lys
		115					120					125			
Val	Phe	Ser	Lys	Glu	Lys	Tyr	Phe	Gln	Thr	He	Lys	Glu	Val	Lys	Glu
	130					135					140				
Ala	Lys	Glu	Lys	Gly	Lys	Lys	Ser	Ser	Arg	Asp	Tyr	۸rg	Arg	Ala	Ala
145					150					155					160
Lys	Tyr	Лsp	Val	Ile	Ser	Val	Gln	Gly	Thr	Glu	Lys	Leu	Ile	Glu	Ala
				165					170					175	
Thr	His	Gly	Glu	Arg	Asp	Arg	Ile	Arg	Tyr	Tyr	Val	His	Lys	Glu	Glu
			180					185					190		
Leu	Phe	Asp	He	Leu	His	Asp	Thr	His	Leu	Ser	Пе	G1y	His	Gly	Gly
		195					200					205			
Arg	Thr	Arg	Met	Leu	Lys	Glu	Leu	Gln	Gly	Lys	Tyr	Gly	Asn	Val	Thr
	210					215					220				
Lys	Glu	Val	lle	Val	Leu	Tyr	Leu	Thr	Leu	Cys	Lys	Gln	Cys	His	Gln
225					230					235					240
Lys	Asn	Pro	Val	Pro	Lys	Arg	Gly	Leu	Ala	Pro	Lys	Pro	Met	Thr	Phe
				245					250					255	
Lys	Asp	lle	Asp	Ser	Thr	Cys	Gln	Val	Glu	lle	Leu	Asp	Met	Gln	Ser
			260					265					270		
Ser	Ala	Asp	Gly	Glu	Phe	Lys	Phe	lle	Leu	Tyr	Tyr	G1n	Asp	His	Ser
		275					280					285			
Thr	Lys	Phe	lle	He	Leu	Arg	Pro	Leu	Arg	Thr	Lys	Gln	Ala	His	Glu
	290					295					300				
Val	Val	Ser	Val	Leu	Leu	Asp	lle	Phe	Thr	lle	Leu	Gly	Thr	Pro	
305					310					315					320
Val	Leu	Asp	Ser	Asp	Ser	Gly	Va]	Glu	Phe	Thr	Asn	Gln	Val	Val	His
				325					330					335	
Glu	Leu	Asn	Glu	Leu	Trp	Pro	Asp	Leu	Lys	lle	Val	Ser	Gly	Lys	Tyr
			340					345					350		
His	Pro	Gly	Gln	Ser	Gln	Gly	Ser	Leu	Glu	Gly	Ala	Ser	Arg	Asp	Val
		355					360					365			
Lys	Asn	Met	11e	Ser	Thr	Trp	Met	Gln	Ser	Asn	His	Ser	Cys	His	Trp
	370					375					380				

Ala Lys Gly Leu Arg Phe Met Gln Met Val Arg Asn Gln Ala Phe Asp Val Ser Leu Gln Gln Ser Pro Phe Glu Ala Met Phe Gly Tyr Lys Ala Lys Phe Gly Leu Tyr Ser Ser Asn Leu Pro Arg Glu Thr Val Ala Thr Leu Gln Thr Glu Glu Glu Leu Glu lle Ala Glu Glu Gln Leu Glu Asn Ser Leu Trp lle Arg Gln Glu Glu Arg Ala Glu Ile Gly Ala Asp Arg Ser Asp Met Asp Asp Asp Met Asp Pro Thr Pro Glu Ala Ser Glu Pro Ser Thr Ser Gln Gly Thr Ser Gly Leu Leu Cys Trp

<210> 4685

<211> 287

<212> PRT

<213> Homo sapiens

<400> 4685

Met Asp Arg Pro Asp Glu Gly Pro Pro Ala Lys Thr Arg Arg Leu Ser Ser Ser Glu Ser Pro Gln Arg Asp Pro Pro Pro Pro Pro Pro Pro Pro Pro Leu Leu Arg Leu Pro Leu Pro Pro Pro Gln Gln Arg Pro Arg Leu Gln Glu Glu Thr Glu Ala Ala Gln Val Leu Ala Asp Met Arg Gly Val Gly Leu Gly Pro Ala Leu Pro Pro Pro Pro Pro Tyr Val lle Leu Glu Glu Gly Gly lle Arg Ala Tyr Phe Thr Leu Gly Ala Glu Cys Pro Gly Trp Asp Ser Thr lle Glu Ser Gly Tyr Gly Glu Ala Pro Pro Pro Thr

Glu Ser Leu Glu Ala Leu Pro Thr Pro Glu Ala Ser Gly Gly Ser Leu Glu lle Asp Phe Gln Val Val Gln Ser Ser Ser Phe Gly Gly Glu Gly Ala Leu Glu Thr Cys Ser Ala Val Gly Trp Ala Pro Gln Arg Leu Val Asp Pro Lys Ser Lys Glu Glu Ala Ile Ile Ile Val Glu Asp Glu Asp Glu Asp Glu Arg Glu Ser Met Arg Ser Ser Arg Arg Arg Arg Arg Arg Arg Arg Arg Lys Gln Arg Lys Val Lys Arg Glu Ser Arg Glu Arg Asn Ala Glu Arg Met Glu Ser IIe Leu Gln Ala Leu Glu Asp IIe Gln Leu Asp Leu Glu Ala Val Asn lle Lys Ala Gly Lys Ala Phe Leu Arg Leu Lys Arg Lys Phe Ile Gln Met Arg Arg Pro Phe Leu Glu Arg Arg Asp Leu Ile Ile Gln His Ile Pro Gly Phe Trp Val Lys Ala Leu His Ser Ser Val Pro Gln Pro Pro Gln Asn Phe Asn Phe Asp Gln Pro Thr

<210> 4686

<211> 196

<212> PRT

<213> Homo sapiens

<400> 4686

 Met Ser Glu Gly Pro Gly Cys Met Gly Ser Glu Gly Gln Leu Cys Pro I
 5
 10
 15

 Trp Ser Glu Gly Arg Gln Cys Pro Arg Ser Glu Gly Cys Gln Cys Ala 20
 25
 30

 Gly Ser Glu Gly Arg His Val Leu Gly Ser Glu Gly Arg Gly Cys Ser 35
 40
 45

Gly Ser Glu Gly Cys Gln Cys Pro Gly Ser Glu Gly Arg Arg Val Leu 55 Arg Phe Glu Gly Trp His Val Leu Gly Ser Glu Gly Gln Arg Val Pro 65 70 75 80 Gly Ser Glu Gly Arg Pro Leu Gly Pro Leu Gly Leu Leu Arg Glu Gly 90 Leu Pro Asp Pro Trp Leu Glu His Arg Pro Arg Gly Asp Pro Ser Pro 105 Ser Gly Ala Leu Pro Arg Pro Pro Ser Leu Ser Phe Leu Thr Trp Phe 115 120 125 Leu Pro Arg Ser Pro Cys Ile Leu Thr Pro Gly Gly Gly Pro Thr Ser 135 140 Asn Ser Leu Ser Pro Ala Gly Ala Trp Ala Gly Pro Arg Ser Leu Leu 150 155 160 145 Pro Gln Leu Ala Phe Arg Gly Glu Thr Lys Ala Gln Gly Leu Phe Trp 165 170 Phe Ser Ala Gln Thr Trp Gln Leu Pro Gly Gly Gly Arg Arg Ala Pro 190 185 Glu Val Gly Ile 195

<210> 4687

<211> 295

<212> PRT

<213> Homo sapiens

<400> 4687

Met Arg Gly Ser Arg Met Ser Gln Pro Pro Gln Cys Leu Arg Ala 1 5 10 15

Gln Ser Ser Cys Cys His Phe Met Val Lys Leu Leu Asp Asp Gly Thr
20 25 30

Phe Met Ile Pro Gly Glu Lys Val Ala His Thr Scr Leu Asp Ala Leu 35 40 45

Val Thr Phe His Gln Gln Lys Pro Ile Glu Pro Arg Arg Glu Leu Leu 50 55 60

Thr	Gln	Pro	Cys	Arg	Gln	Lys	Asp	Pro	Ala	Asn	Val	Asp	Tyr	Glu	Asp
65					70					75					80
Leu	Phe	Leu	Tyr	Ser	Asn	Ala	Va]	Ala	Glu	Glu	Ala	Ala	Cys	Pro	Val
				85					90					95	
Ser	Ala	Pro	Glu	Glu	Ala	Ser	Pro	Lys	Pro	Val	Leu	Cys	His	Gln	Ser
			100					105					110		
Lys	Glu	Arg	Lys	Pro	Ser	Ala	Glu	Met	Asn	Gly	He	Thr	Thr	Lys	G] u
		115					120					125			
Ala	Thr	Ser	Ser	Cys	Pro	Pro	Lys	Ser	Pro	Leu	Gly	${\sf Glu}$	Thr	Arg	Gln
	130					135					140				
Lys	Leu	Trp	Arg	Ser	Leu	Lys	Met	Leu	Pro	Glu	Arg	Gly	Gln	Arg	Val
145					150					155					160
Arg	Gln	Gln	Leu	Lys	Ser	His	Leu	Ala	Thr	Val	Asn	Leu	Ser	Ser	Leu
				165					170					175	
Leu	Asp	Val	Arg	Arg	Ser	Thr	Val	Пе	Ser	Gly	Pro	Gly	Thr	Gly	Lys
			180					185					190		
Gly	Ser	Gln	Asp	His	Ser	Gly	Asp	Pro	Thr	Ser	Gly	Asp	Arg	Gly	Tyr
		195					200					205			
Thr	Asp	Pro	Cys	Val	Ala	Thr	Ser	Leu	Lys	Ser	Pro	Ser	Gln	Pro	Gln
	210					215					220				
Ala	Pro	Lys	Asp	Arg	Lys	Val	Pro	Thr	Arg	Lys	Ala	Glu	Arg	Ser	Va]
225					230					235					240
Ser	Cys	He	Glu	Val	Thr	Pro	Gly	Asp	Arg	Ser	Trp	His	Gln	Met	Val
				245					250					255	
Val	Arg	Ala	Leu	Ser	Ser	Gln	Glu	Ser	Lys	Pro	Glu	His	Gln	Gly	Leu
			260					265					270		
Ala	Glu	Pro	Glu	Asn	Asp	Gln	Leu	Pro	Glu	Glu	Tyr	Gln	Gln	Pro	Pro
		275					280					285			
Pro	Phe	Ala	Pro	Gly	Tyr	Cys									
	290					295									

<211> 594

<212> PRT

<213> Homo sapiens

<400)> 46	888													
Met	Lys	Val	Thr	Leu	Ser	Ala	Leu	Asp	Thr	Ser	Glu	Ser	Ser	Phe	Thr
1				5					10					15	
Pro	Leu	Val	Val	lle	Glu	Leu	Ala	Gln	Asp	Val	Lys	Glu	Glu	Thr	Lys
			20					25					30		
Glu	Trp	Leu	Lys	Asn	Arg	He	11e	Ala	Lys	Lys	Lys	Лѕр	Gly	Asp	Asn
		35					40					45			
Asn	Asp	Asp	Phe	Leu	Thr	Met	Ala	Glu	Cys	Gln	Phe	Ile	Ile	Lys	His
	50					55					60				
Glu	Leu	Glu	Asn	Leu	Arg	Ala	Lys	Asp	Glu	Lys	Met	Ile	Pro	Gly	Tyr
65					70					75					80
Pro	Gln	Ala	Lys	Leu	Tyr	Pro	Gly	Lys	Ser	Leu	Leu	Arg	Arg	Leu	Leu
				85					90					95	
Thr	Ser	Gly	lle	Val	He	Gln	Val	Phe	Pro	Leu	His	Asp	Ser	Glu	Ala
			100					105					110		
Leu	Lys	Lys	Leu	Glu	Asp	Thr	Trp	Tyr	Thr	Arg	Phe	Ala	Leu	Lys	Tyr
		115					120					125			
Gln	Pro	11e	Asp	Ser	Ile	Arg	Gly	Tyr	Phe	Gly	Glu	Thr	lle	Ala	Leu
	130					135					140				
Tyr	Phe	Gly	Phe	Leu	Glu	Tyr	Phe	Thr	Phe	Ala	Leu	lle	Pro	Met	Ala
145					150					155					160
Val	lle	Gly	Leu	Pro	Tyr	Туг	Leu	Phe	Val	Trp	Glu	Asp	Tyr	Asp	Lys
				165					170					175	
Tyr	Val	lle	Phe	Ala	Ser	Phe	Asn	Leu	He	Trp	Ser	Thr	Val	Пе	Leu
			180					185					190		
Glu	Leu	Trp	Lys	Arg	Gly	Cys	Ala	Asn	Met	Thr	Tyr	Arg	Trp	Gly	Thr
		195					200					205			
Leu	Leu	Met	Lys	Arg	Lys	Phe	Glu	Glu	Pro	Arg	Pro	Gly	Phe	His	Gly
	210					215					220				
Val	Leu	Gly	Ile	Asn	Ser	lle	Thr	Gly	Lys		Glu	Pro	Leu	Tyr	Pro
225					230					235					240
Ser	Tyr	Lys	Arg		Leu	Arg	He	Tyr		Val	Ser	Leu	Pro		Va]
				245					250					255	
Cys	Leu	Cys		Tyr	Phe	Ser	Leu		Val	Met	Met	He		Phe	Asp
			260					265					270		

Met	Glu	Val	Trp	Ala	Leu	Gly	Leu	His	Glu	Asn	Ser	Gly	Ser	Glu	Trp
		275					280					285			
Thr	Ser	Val	Leu	Leu	Tyr	Val	Pro	Ser	lle	He	Tyr	Ala	He	Val	lle
	290					295					300				
Glu	11e	Met	Asn	Arg	Leu	Tyr	Arg	Tyr	Ala	Ala	Glu	Phe	Leu	Thr	Ser
305					310					315					320
Trp	Glu	Asn	His	Arg	Leu	Glu	Ser	Ala	Tyr	Gln	Asn	His	Leu	lle	Leu
				325					330					335	
Lys	Val	Leu	Val	Phe	Asn	Phe	Leu	Asn	Cys	Phe	Ala	Ser	Leu	Phe	Tyr
			340					345					350		
Ile	Ala	Phe	Val	Leu	Lys	Asp	Met	Lys	Leu	Leu	Arg	Gln	Ser	Leu	Ala
		355					360					365			
Thr	Leu	Leu	He	Thr	Ser	Gln	11e	Leu	Asn	Gln	He	Met	Glu	Ser	Phe
	370					375					380				
Leu	Pro	Tyr	Trp	Leu	Gln	Arg	Lys	His	Gly	Val	Gln	Val	Lys	Arg	Lys
385					390					395					400
Val	Gln	Ala	Leu	Lys	Ala	Asp	Ile	Asp	Ala	Thr	Leu	Tyr	Glu	Gln	Val
				405					410					415	
Ile	Leu	Glu	Lys	Glu	Met	Gly	Thr	Tyr	Leu	Gly	Thr	Phe	Asp	Asp	Tyr
			420					425					430		
Leu	Glu	Leu	Phe	Leů	Gln	Phe	Gly	Tyr	Val	Ser	Leu	Phe	Ser	Cys	Val
		435					440					445			
Tyr	Pro	Leu	Ala	Ala	Ala	Phe	Ala	Val	Leu	Asn	Asn	Phe	Thr	Glu	Val
	450					455					460				
	Ser	Asp	Ala	Leu		Met	Cys	Arg	Val		Lys	Arg	Pro	Phe	
465					470					475					480
Glu	Pro	Ser	Ala		lle	Gly	Val	Trp		Leu	Ala	Phe	Glu		Met
_				485					490					495	
Ser	Val	He		Val	Val	Thr	Asn		Ala	Leu	He	G1 y		Ser	Pro
			500		ъ.			505			_		510		
Gln	Val		Ala	Val	Phe	Pro		Ser	Lys	Ala	Asp		He	Leu	He
	17 1	515	V 1	C1			520	,				525 B			
val		Ala	val	Glu	HIS		Leu	Leu	Ala	Leu	Lys	Phe	11e	Leu	Ala
DI	530	11.	D	Λ - · ·		535	Α	11.2	11	C)	540			A 7	Δ.
	Ala	116	rro	лѕр		Pro	Arg	H1S	116		Met	Lys	Leu	Ala	
545					550					555					560

Leu Glu Phe Glu Ser Leu Glu Ala Leu Lys Gln Gln Gln Met Lys Leu 565 570 575 575 Val Thr Glu Asn Leu Lys Glu Glu Pro Met Glu Ser Gly Lys Glu Lys 580 585 585 590

Ala Thr

<210> 4689

<211> 120

<212> PRT

<213> Homo sapiens

<400> 4689

Met Phe Leu Ala Pro Leu Leu His Ser Leu Val Leu Pro Gly Leu Phe
1 5 10 15

Leu Ala Pro Pro Leu His Ser Ala Val Leu Pro Gly Met Phe Leu Ala 20 25 30

Pro Pro Leu His Ser Pro Val Leu Pro Gly Met Phe Leu Ala Pro Pro 35 40 45

Leu His Ser Pro Val Leu Ile Ser His Phe Ser Lys Val Gly Phe Arg 50 55 60

Gly Arg Arg Asp Glu Arg Lys Asp Thr Ala Ser His Gly Gly Tyr Gly
65 70 75 80

Ser His Leu Val Met Gly Leu Ser Gly Cys Asp Lys Tyr Thr Lys Pro 85 90 95

His Phe Phe Ser Trp Gly Ala Phe Gly Glu Leu Leu Trp Arg Pro Ser 100 105 110

His Arg Met Leu Glu Asp Gly Phe 115 120

<210> 4690

<211> 461

<212> PRT

<213> Homo sapiens

<400)> 46	590													
Met	Glu	Arg	Glu	Gly	He	Trp	His	Ser	Thr	Leu	Gly	Glu	Thr	Trp	Glu
1				5					10					15	
Pro	Asn	Asn	Trp	Leu	Glu	Gly	Gln	Gln	Asp	Ser	His	Leu	Ser	Gln	Val
			.20					25					30		
Gly	Val	Thr	His	Lys	Glu	Thr	Phe	Thr	Glu	Met	Arg	Val	Cys	Gly	Gly
		35					40					45			
Asn	Glu	Phe	Glu	Arg	Cys	Ser	Ser	Gln	Asp	Ser	Ile	Leu	Asp	Thr	Gln
	50					55					60				
Gln	Ser	Ile	Pro	Met	Val	Lys	Arg	Pro	His	Asn	Cys	Asn	Ser	His	Gly
65					70					75					80
Glu	Asp	Ala	Thr	G]n	Asn	Ser	Glu	Leu	lle	Lys	Thr	Gln	Arg	Met	Phe
				85					90					95	
Val	Gly	Lys	Lys	lle	Tyr	Glu	Cys		Gln	Cys	Ser	Lys	Thr	Phe	Ser
			100					105					110		
Gln	Ser	Ser	Ser	Leu	Leu	Lys		Gln	Arg	Ile	His		Gly	Glu	Lys
		115					120					125		_	_
Pro		Lys	Cys	Asn	Val		Gly	Lys	His	Phe		Glu	Arg	Ser	Ser
	130			0.1		135		m.	0.1	0.1	140		m		
	lhr	Val	His	GIn		He	HIS	Ihr	Gly		Lys	Pro	lyr	Lys	
145	C1	C	C1	1	150	DI	C .	C1	C .	155	4		TI.	W 3	160
Asn	GIU	Cys	GIY		АТа	rne	ser	61n		мет	Asn	Leu	inr		HIS
Cla	A 22.00	Thn	uio	165	Clu	Clu	lva	Dro	170	Cln	Cva	Luc	C1.,	175	Cly
GIII	A1 g	Thr	180	1111	Gly	Gju	LyS	185	1 y 1	OIII	Cys	LyS	190	Cys	оту
Lvc	Δla	Phe		Lve	Aen	Sor	Sor		ماآ	Gln	Hic	Glu		ماا	Hic
LyS	ліа	195		Lys	ЛЗП	361		Leu		Oin	1115	205		116	1113
Thr	Glv	Glu		Pro	Tvr	Lvs				Cvs	Glv			Phe	Thr
	210	014	12 3 13	,,,	.,1	215	0,0	71011	0.4	0,0	220	Ljo	7170	1110	
Gln		Met	Asn	Leu	Thr		His	Gln	Arg	Thr		Thr	Glv	Glu	Lvs
225				.500	230				0	235		••••	01,	0.2.0	240
	Tyr	Glu	Cvs	Asn		Cvs	Gly	Lvs	Ala		Ser	Gln	Ser	Met	
	y -		•	245		•	- •	•	250					255	
Leu	lle	Val	His		Arg	Ser	His	Thr		Glu	Lys	Pro	Tyr		Cys
			260					265					270		

Ser Gln Cys Gly Lys Ala Phe Ser Lys Ser Ser Thr Leu Thr Leu His Gln Arg Asn His Thr Gly Glu Lys Pro Tyr Lys Cys Asn Lys Cys Gly Lys Ser Phe Ser Gln Ser Thr Tyr Leu Ile Glu His Gln Arg Leu His Ser Gly Val Lys Pro Phe Glu Cys Asn Glu Cys Gly Lys Ala Phe Ser Lys Asn Ser Ser Leu Thr Gln His Arg Arg Ile His Thr Gly Glu Lys Pro Tyr Glu Cys Met Val Cys Gly Lys His Phe Thr Gly Arg Ser Ser Leu Thr Val His Gln Val lle His Thr Gly Glu Lys Pro Tyr Glu Cys Asn Glu Cys Gly Lys Ala Phe Ser Gln Ser Ala Tyr Leu Ile Glu His Gln Arg Ile His Thr Gly Glu Lys Pro Tyr Glu Cys Asp Gln Cys Gly Lys Ala Phe Ile Lys Asn Ser Ser Leu Thr Val His Gln Arg Thr His Thr Gly Glu Lys Pro Tyr Gln Cys Asn Glu Cys Gly Lys Ala Phe Ser Arg Ser Thr Asn Leu Thr Arg His Gln Arg Thr His Thr

<210> 4691

<211> 122

<212> PRT

<213> Homo sapiens

<400> 4691

Met Thr 11e Ala Thr Pro Phe Val Cys Leu Ile Phe Ala His Ser Gln 1 5 10 15 Val Phe Thr Asp Phe 11e Pro Tyr Ser Arg His Leu Gln Met Pro Thr 20 25 30

 Ser
 Tyr
 Pro
 Cys
 Ser
 Lys
 Thr
 His
 Phe
 Lys
 Cys
 Pro
 Leu
 Leu
 Trp
 Glu

 Met
 Pro
 Val
 Val
 Pro
 Leu
 Asp
 Thr
 Asn
 His
 Leu
 Ser
 Ser
 Leu
 Glu
 Pro

 50
 Thr
 His
 Ala
 Pro
 Thr
 Pro
 Ile
 Ser
 Asn
 Ala
 Tyr
 Ser
 Ser
 Arg
 Lys

 65
 Tyr
 Fro
 Thr
 His
 Gly
 Gln
 Lys
 Ser
 Leu
 Leu
 Leu
 Thr
 Ala
 Asn
 Asn

 Cys
 Pro
 Ser
 Thr
 His
 Gly
 Gln
 Lys
 Ser
 Leu
 Leu
 Leu
 Thr
 Ala
 Asn

 Cys
 Pro
 Ser
 Thr
 His
 Leu
 Ser
 His
 Val
 Leu
 Ser
 Ile
 Thr

 Lys
 Tyr
 Pro
 His
 Leu
 Leu
 Leu
 Leu
 Thr

<210> 4692

<211> 123

<212> PRT

<213> Homo sapiens

⟨400⟩ 4692

Met Lys Gly Pro Cys Pro His Leu Pro Pro Pro Leu Ser Ser Cys Ile
1 5 10 15

Met Asn Glu Thr Ala Ala Ser Leu Leu Pro Glu Val Leu His Phe Arg 20 25 30

Leu Gly Cys Asn Gly Ser 11e Ser Ala Gln Cys Asn Leu Cys Phe Pro 35 40 45

Gly Ser Ser Asp Ser Pro Ala Ser Ala Ser Gln Ala Ala Val Asn Thr 50 55 60

Gly Trp Ser Ala Val Val Leu Cys Leu Glu Phe Val Pro Ala Val Gly
65 70 75 80

Phe Val Val Leu Leu Thr Ser Arg Met Lys Pro Arg Thr Phe Thr Arg

85 90 95

Asn Ile Thr Leu Tyr Gly Cys Thr Thr Val Cys Leu Ser Ile Leu Gln 100 105 110

Val Lys Asp Met Trp Val Val Pro Ser Ala Gly

<210> 4693 <211> 127 <212> PRT <213> Homo sapiens <400> 4693 Met Asp Leu His His Ser Val Gln Pro Glu Ser Leu Leu Leu Ser Thr 1 5 10 15 Ala Ser Phe Ser Phe Ser Leu Ile Ile Pro Phe Asn Ser Asn Lys Met 25 Thr Trp Pro Ala His Glu Asn Thr Glu Met Arg Pro His Leu Val Ile 35 45 40 Ser Gly Val Leu Arg Gly Thr Leu Val Val Leu Gly Ala Ala Val 55 60 Leu Ser Val Lys Asn Phe Gln Glu Phe Leu lle Ser Cys Phe His Gln 70 75 80 Asp Ser His Asn Leu Leu Leu Leu Pro Leu Ser Ser Gly Phe Val Pro 90 85 Glu His lle Ile Arg Lys Ala Ala Ile Ile Thr Ala Tyr Leu Pro Pro 105 Ala Pro Leu His Lys His Pro Pro Ser Pro His Cys Ala Lys Gln 115 120 125 <210> 4694 <211> 194 <212> PRT <213> Homo sapiens

1 5 10 15

Ser Ala Ser Ser Gly Asn Phe Ile Pro Pro Tyr His Pro Ser Ser Leu
20 25 30

Met Asp Arg Pro Ser Leu Val Arg Ser Arg Lys Gln Ser Pro Arg Leu

<400> 4694

Gly Lys Val Glu Arg Thr Asn Gly Leu Leu Lys Thr His Leu Thr Lys 40 Leu Ser Leu Gln Leu Lys Lys Asp Trp Thr Ala Leu Leu Pro Phe Ala 50 55 60 Leu Leu Arg Ile Arg Ala Tyr Pro Gln Glu Ala Thr Gly Tyr Ser Pro 70 75 Phe Glu Leu Ser Tyr Gly Cys Thr Phe Leu Leu Gly Pro Asn Leu Leu 85 90 Thr Asp Asn Thr Tyr Ala Asp Met Gln Gln Lys Lys Gln Leu Val Phe 100 105 110 Pro His Leu Ser Leu Thr Ala Ser Phe Leu Pro Ser His Leu Ser Leu 120 Pro Pro Ile Leu Pro Leu Lys Leu Leu Pro Ile Asn Pro Phe Leu Leu 130 135 140 Met Lys Leu Leu Pro 11e Asn Pro Phe Leu Leu Lys Ala Asn Gly Ser 145 150 155 160 Trp Ile Lys Glu Asn Ser Ser Phe Leu Pro His Arg Leu Ile Leu Ser 165 170 Tyr His Pro Phe Arg Thr Ser Phe Met Trp Val Thr Ser Gln Trp Pro 180 185 190 lle Ser

<210> 4695

<211> 237

<212> PRT

<213> Homo sapiens

<400> 4695

Met Ser Cys Arg His Ser Phe Pro Leu Leu Pro Cys Ser Leu His

1 5 10 15

Ser His Ala Phe Leu Ser Ser Cys His Gly Asp Leu Cys Pro Leu Thr
20 25 30

Ser His Thr Ala Cys Phe Leu Ala Cys Val Pro Pro Gln Ala Gly Gly

		35					40					45			
His	Phe	Gln	Ser	Ser	Phe	Thr	Pro	Cys	Val	Arg	Val	He	Asn	Gln	He
	50					55					60				
Trp	Ala	Pro	Val	Leu	His	Pro	Leu	Cys	His	Asn	Ser	Pro	Cys	Pro	Pro
65					70					75					80
Asn	Leu	Pro	Gly	Thr	Arg	Phe	Gly	Ala	Met	Leu	Asp	Met	Leu	Thr	Asp
				85					90					95	
Arg	Cys	Ser	Thr	Met	Cys	Leu	Leu	Val	Asn	Leu	Ala	Leu	Leu	Tyr	Pro
			100					105					110		
Gly	Ala	Thr	Leu	Phe	Phe	Gln	Ile	Ser	Met	Ser	Leu	Asp	Val	Ala	Ser
		115					120					125			
His	Trp	Leu	His	Leu	His	Ser	Ser	Val	Val	Arg	Gly	Ser	Glu	Ser	His
	130					135					140				
Lys	Met	lle	Asp	Leu	Ser	Gly	Asn	Pro	Val	Leu	Arg	lle	Tyr	Tyr	Thr
145					150					155					160
Ser	Arg	Pro	Ala	Leu	Phe	Thr	Leu	Cys	Ala	Gly	Asn	Glu	Leu	Phe	Tyr
				165					170					175	
Cys	Leu	Leu	Tyr	Leu	Phe	His	Phe	Ser	Glu	Gly	Pro	Leu	Val	Gly	Ser
			180					185					190		
Va]	Gly	Leu	Phe	Arg	Met	Gly	Leu	Trp	Val	Thr	Ala	Pro	lle	Ala	Leu
		195					200					205			
Leu	Lys	Ser	Leu	Ile	Ser	Val	Ile	His	Leu	11e	Thr	Ala	Ala	Arg	Asn
	210					215					220				
Met	Ala	Ala	Leu	Asp	Ala	Ala	Asp	Arg	Ala	Lys	Lys	Lys			
225					230					235					

<211> 111

<212> PRT

<213> Homo sapiens

<400> 4696

			20					25					30		
Leu	Pro	Asp	Leu	Leu	Leu	Pro	Gly	Pro	Ser	Gln	Ala	Tyr	Pro	Phe	Leu
		35					40					45			
Ser	Ser	Leu	Pro	Ser	Phe	Thr	Pro	Ser	Leu	Ser	Met	Glu	Pro	His	Ser
	50					55					60				
Gly	GIn	Λsp	His	Leu	Leu	Trp	Pro	Ser	Glu	Gly	Phe	Arg	Thr	Ala	Gly
65					70					75					80
Gly	Cys	Ser	Pro	Arg	Pro	Leu	Glu	Trp	Phe	Val	lle	Phe	Leu	Cys	Leu
				85					90					95	
Val	Pro	Thr	Cys	Leu	Ser	Ser	Leu	Leu	Pro	Pro	His	Ala	Ser	Ser	
			100					105					110		
<21	0> 46	597													
	1> 13														
	2> PF														
<21	3> Ho	omo :	sapi	ens											
	0> 46		,			14 .		,	17.1	41.	Δ	C1	1	C1	C1
	Asn	Glu	Lys		Lys	Met	Leu	Lys		Ala	Arg	Glu	Lys	15	GIR
1	Tl	Т	Lua	5	Luc	Dno	Tlo	Ana	10	Mot	Val	C1v	Lou		Ala
val	Thr	ıyr	Lys 20	GIY	Lys	PTO	116	71 g 25	reu	Me t	vai	GIY	30	Leu	nia
Aan	Thr	Lou		Ala	110	Ana	Acn		C1v	Pro	Mot	Pho		مات	Sor
ASP	1111	35	OIII	MIA	116	ΛI g	40	Leu	Oly	1.10	MC C	45	ASII	116	561
Phe	Leu		Phe	Ser	Phe	Leu		Phe	Phe	Leu	Asn		Val	Ser	l.eu
1110	50	1110	The	561	1 110	55	1110	1110	1110		60	6			24-
Cvs	llis	Gln	Ala	Glv	Val		Trp	His	Asp	Leu		Ser	Leu	Gln	Pro
65		0111		0,,	70		4			75					80
	Pro	Pro	Arg	Phe		Gln	Phe	Ser	Cvs			Leu	Pro	Ser	Ser
				85					90					95	
Trn	Asp	Tyr	Arg	85 Arg	Ala	Pro	Ser	His			Asn	Phe	Cys		Phe
Trp	Asp	Tyr	Arg 100		Ala	Pro	Ser	His	Leu		Asn	Phe	Cys		Phe
	Asp Arg		100	Arg				105	Leu	Ala			110	lle	

Asp Phe Met 11e

<210> 4698 <211> 665 <212> PRT <213> Homo sapiens <400> 4698 Met Met Lys Thr Phe Phe Ser Thr Gly Gln Gly Asp Thr Glu Ala Phe His Thr Gly Thr Leu Gln Arg Gln Ala Ser His His 11e Gly Asp Phe Cys Phe Gln Lys lle Glu Lys Asp lle His Gly Phe Gln Phe Gln Trp Lys Glu Asp Glu Thr Asn Asp His Ala Ala Pro Met Thr Glu lle Lys Glu Leu Thr Gly Ser Thr Gly Gln His Asp Gln Arg His Ala Gly Asn Lys His 11e Lys Asp Gln Leu Gly Leu Ser Phe His Ser His Leu Pro Glu Leu His 11e Phe Gln Pro Glu Gly Lys 11e Gly Asn Gln Val Glu Lys Ser Ile Asn Asn Ala Ser Ser Val Ser Thr Ser Gln Arg Ile Cys Cys Arg Pro Lys Thr His lle Ser Asn Lys Tyr Gly Asn Asn Ser Leu His Ser Ser Leu Leu Thr Gln Lys Arg Asn Val His Met Arg Glu Lys Ser Phe Gln Cys Ile Glu Ser Gly Lys Ser Phe Asn Cys Ser Ser Leu Leu Lys Lys His Gln 11e Thr His Leu Glu Glu Lys Gln Cys Lys Cys Asp Val Tyr Gly Lys Val Phe Asn Gln Lys Arg Tyr Leu Ala Cys His

Arg	Arg	Ser	His	lle	Asp	Glu	Lys	Pro	Tyr	Lys	Cys	Asn	Glu	Cys	Gly
	210					215					220				
Lys]]e	Phe	Gly	His	Asn	Thr	Ser	Leu	Phe	Leu	His	Lys	Ala	Leu	His
225					230					235					240
Thr	Ala	Asp	Lys	Pro	Tyr	Glu	Cys	Glu	Glu	Cys	Asp	Lys	Val	Phe	Ser
				245					250					255	
Arg	Lys	Ser	His	Leu	Glu	Thr	His	Lys	He	He	Tyr	Thr	Gly	Gly	Lys
			260					265					270		
Pro	Tyr	Lys	Cys	Lys	Val	Cys	Asp	Lys	Ala	Phe	Thr	Cys	Asn	Ser	Tyr
		275					280					285			
Leu	Ala	Lys	His	Thr	Ile	He	His	Thr	Gly	Glu	Lys	Pro	Tyr	Lys	Cys
	290					295					300				
Asn	Glu	Cys	Gly	Lys	Va]	Phe	Asn	Arg	Leu	Ser	Thr	Leu	Ala	Arg	His
305					310					315					320
Arg	Arg	Leu	His	Thr	Gly	Glu	Lys	Pro	Tyr	Glu	Cys	Glu	Glu	Cys	Glu
				325					330					335	
Lys	Val	Phe	Ser	Arg	Lys	Ser	His	Leu	Glu	Arg	His	Lys	Arg	He	His
			340					345					350		
Thr	Gly	Glu	Lys	Pro	Tyr	Lys	Cys	Lys	Val	Cys	Asp	Lys	Ala	Phe	Ala
		355					360					365			
Tyr	Asn	Ser	Tyr	Leu	Ala	Lys	His	Ser	He	lle	His	Thr	Gly	Glu	Lys
	370					375					380				
	Tyr	Lys	Cys	Asn	Glu	Cys	Gly	Lys	Val		Asn	Gln	Gln	Ser	
385					390					395					400
Leu	Ala	Arg	His		Arg	Leu	His	Thr		Glu	Gln	Pro	Tyr		Cys
				405					410					415	
G] u	Glu	Cys		Lys	Va1	Phe	Arg		Lys	Ser	His	Leu		Arg	His
			420				_	425					430		
Arg	Arg		His	Thr	Gly	G] u		Pro	Tyr	Lys	Cys		Val	Cys	Asp
		435					440					445			
Lys		Phe	Arg	Ser	Asp	Ser	Cys	Leu	Thr	Glu		GIn	Arg	Val	His
	450					455					460				
	Gly	Glu	Lys	Pro		Thr	Cys	Asn	GJu		GIV	Lys	Val	Phe	
465	,	4.1			470		11.			475	12.	Tr.	A 1	61	480
Thr	Lys	Ala	Asn		Ala	Cys	His	His		Leu	HIS	Thr	Λla		Lys
				485					490					495	

Pro Tyr Lys Cys Glu Glu Cys Glu Lys Val Phe Ser Arg Lys Ser His Met Glu Arg His Arg Arg Ile His Thr Gly Glu Lys Pro Tyr Lys Cys Lys Val Cys Asp Lys Ala Phe Arg Arg Asp Ser His Leu Ala Gln His Gln Arg Val His Thr Gly Glu Lys Pro Tyr Lys Cys Asn Glu Cys Gly Lys Thr Phe Arg Gln Thr Ser Ser Leu lle Ile His Arg Arg Leu His Thr Gly Glu Lys Pro Tyr Lys Cys Asn Glu Cys Gly Lys Thr Phe Ser Gln Met Ser Ser Leu Val Tyr His His Arg Leu His Ser Gly Glu Lys Pro Tyr Lys Cys Asn Glu Cys Gly Lys Val Phe Asn Gln Gln Ala His Leu Ala Gln His Gln Arg Val His Thr Gly Glu Lys Pro Tyr Lys Cys Asn Glu Cys Gly Lys Thr Phe Ser Gln Met Ser Asn Leu Val Tyr His His Arg Leu His Ser Gly Glu Lys Pro

<210> 4699

<211> 986

<212> PRT

<213> Homo sapiens

<400> 4699

Met Pro Arg Ala Thr Trp Ala Asn Ser Lys Glu Arg Ser Trp Ala Glu 1 5 10 15 Ser Glu Arg Gly Pro Arg Asp Thr Gly Asn Gly Gly Ser Lys Ala Glu 20 25 30 Arg His Ile Gln Glu Ile Glu Thr Gly Arg Gly Gly Asp Arg Ala Lys 35 40 45

Ala		Arg	Arg	Gln	Arg	Met	Arg	Leu	Arg	Gly	Thr 60	Glu	Arg	Ala	Ser
	50	D	C1	Δ	A	55	C1	A a.u.	Com	A 22 cr		The	Aan	Mot	Duo
	бту	Pro	оту	Arg		Leu	GIY	ASP	261.		Gry	1 (1.3	ASP	wet	
65	4.3		4.1	C1	70	1		A 1 -	۸1	75 Mat	Than	C1	C1	Can	80
бту	Ala	Arg	Ala		оту	Leu	ATA	мта		me t	1111.	GIU	Gru		010
61	T)	17 1	,	85 T	7.1	C1	11.2	A	90 T	V . 1	Contra	C	C1	95	۸
Glu	Ihr	vai		lyr	11e	Glu	H1 S		ıyr	vai	Cys	ser		Cys	ASI
61	,	T	100	C		C1	C1	105	1	W-4	U	C1	110	C	11: -
GIn	Leu	-	Gly	Ser	Leu	Glu		val	Leu	Met	нıs		Asn	ser	HIS
., 1	Б	115	61		D1	61	120	17 7	01	v 3	4.3	125	D	C1	v 1
Val		Gln	GIn	HIS	Phe	Glu	Leu	val	Gly	vai		Asp	Pro	GIY	vai
	130		m)		mı	135		0.1	m.	61	140	Tr.	C.I.	T)	
	Val	Ala	Thr	Asp		Ala	Ser	GIy	lhr		Leu	lyr	GIn	Ihr	
145					150	~1	_	,	0.1	155	61	C 1	,		160
Val	GIn	Glu	Ser		Tyr	Gln	Cys	Leu		Cys	61y	GIn	Leu		Met
~		0	0.1	165		0.1		0.1	170			,	,	175	
Ser	Pro	Ser		Leu	Leu	Glu	His		Glu	Leu	His	Leu		Met	Met
			180			_		185					190		
Ala	Pro		GIu	Ala	Val	Pro		Glu	Pro	Ser	Pro		Ala	Pro	Pro
		195		m.			200	0.1	0	17 1		205	,	4.7	
Leu		Ser	Ser	Thr	He	His	lyr	Glu	Cys	Val		Cys	Lys	Ala	Leu
	210	_				215					220	ar i			
	Ala	Ser	GIn	Glu		Trp	Leu	Asn	His		GIn	lhr	His	Leu	
225		_	 .		230					235		0.1	0	10	240
Ala	Thr	Pro	Thr		Ala	Pro	Ala	Pro		Val	Leu	Gly	Ser		Val
				245			0.1		250				0.1	255	C
Val	Leu	Gly		Pro	Val	Gly	GIn		Arg	Val	Ala	Val		НIS	Ser
_			260	0.1	0.1	0.1	0.1	265	0.1		mı.		270	C	
Tyr	Arg		Ala	Glu	Glu	Gly		GIu	Gly	Ala	Ihr		Pro	Ser	Ala
		275					280		9.3			285			ar.
Ala		Thr	Thr	Thr	Glu	Val	Val	Thr	Glu	Val		Leu	Leu	Leu	lyr
	290	_				295			0.3		300			131	
•	Cys	Ser	Glu	Cys		G1n	Leu	Phe	GIn		Pro	Ala	Asp	Phe	
305		0.3		m.	310	D.				315	15	6.3	6	0.1	320
Glu	His	GIn	Ala		His	Phe	Pro	Mla		Val	Pro	Glu	Ser		ыu
				325					330		•			335	

Pro	Ala	Leu	Gln	Gln	Glu	Val	Gln	Ala	Ser	Ser	Pro	Ala	Glu	Val	Pro
			340					345					350		
Val	Ser	Gln	Pro	Asp	Pro	Leu	Pro	Ala	Ser	Λsp	His	Ser	Tyr	Glu	Leu
		355					360					365			
Arg	Asn	Gly	Glu	Ala	Ile	Gly	Arg	Asp	Arg	Arg	Gly	Arg	Arg	Ala	Arg
	370					375					380				
Arg	Asn	Asn	Ser	Gly	Glu	Ala	Gly	Gly	Ala	Ala	Thr	Gln	Glu	Leu	Phe
385					390					395					400
Cys	Ser	Ala	Cys	Asp	Gln	Leu	Phe	Leu	Ser	Pro	His	Gln	Leu	Gln	Gln
				405					410					415	
His	Leu	Arg	Ser	His	Arg	Glu	Gly	Val	Phe	Lys	Cys	Pro	Leu	Cys	Ser
			420					425					430		
Arg	Val	Phe	Pro	Ser	Pro	Ser	Ser	Leu	Asp	Gln	His	Leu	Gly	Asp	His
		435					440					445			
Ser	Ser	Glu	Ser	His	Phe	Leu	Cys	Val	Asp	Cys	Gly	Leu	Ala	Phe	Gly
	450					455					460				
Thr	Glu	Ala	Leu	Leu	Leu	Ala	His	Arg	Arg	Ala	His	Thr	Pro	Asn	Pro
465					470					475					480
Leu	His	Ser	Cys	Pro	Cys	Gly	Lys	Thr	Phe	Val	Asn	Leu	Thr	Lys	Phe
				485					490					495	
Leu	Tyr	His	Arg	Arg	Thr	His	Gly	Val	Gly	Gly	Val	Pro	Leu	Pro	Thr
			500					505					510		
Thr	Pro	Val	Pro	Pro	Glu	Glu	Pro	Val	lle	Gly	Phe	Pro	Glu	Pro	Ala
		515					520					525			
Pro	Ala	Glu	Thr	Gly	Glu	Pro	Glu	Ala	Pro	Glu	Pro	Pro	Val	Ser	Glu
	530					535					540				
Glu	Thr	Ser	Ala	Gly	Pro	Ala	Ala	Pro	Gly	Thr	Tyr	Arg	Cys	Leu	Leu
545					550					555					560
Cys	Ser	Arg	Glu	Phe	Gly	Lys	Ala	Leu	G]n	Leu	Thr	Arg	His	Gln	Arg
				565					570					575	
Phe	Val	His	Arg	Leu	Glu	Arg	Arg	His	Lys	Cys	Ser	He	Cys	Gly	Lys
			580					585					590		
Met	Phe	Lys	Lys	Lys	Ser	His	Val	Arg	Asn	His	Leu	Arg	Thr	His	Thr
		595					600					605			
Gly	Glu	Arg	Pro	Phe	Pro	Cys	Pro	Asp	Cys	Ser	Lys	Pro	Phe	Asn	Ser
	610					615					620				

Pro	Ala	Asn	Leu	Ala	Arg	His	Arg	Leu	Thr	His	Thr	Gly	Glu	Arg	Pro
625					630					635					640
Tyr	Arg	Cys	Gly	Asp	Cys	Gly	Lys	Ala	Phe	Thr	Gln	Ser	Ser	Thr	Leu
				645					650					655	
Arg	Gln	His	Arg	Leu	Val	His	Ala	Gln	His	Phe	Pro	Tyr	Arg	Cys	Gln
			660					665					670		
Glu	Cys	Gly	Val	Arg	Phe	His	Arg	Pro	Tyr	Arg	Leu	Leu	Met	His	Arg
		675					680					685			
Tyr	His	His	Thr	Gly	Glu	Tyr	Pro	Tyr	Lys	Cys	Arg	Glu	Cys	Pro	Arg
	690					695					700				
Ser	Phe	Leu	Leu	Arg	Arg	Leu	Leu	Glu	Val	His	Gln	Leu	Val	Val	His
705					710					715					720
Ala	Gly	Arg	Gln	Pro	His	Arg	Cys	Pro	Ser	Cys	Gly	Ala	Ala	Phe	Pro
				725					730					735	
Ser	Ser	Leu	Arg	Leu	Arg	Glu	His	Arg	Cys	Ala	Ala	Ala	Ala	Ala	Gln
			740					745					750		
Ala	Pro		Arg	Phe	Glu	Cys		Thr	Cys	Gly	Lys		Val	Gly	Ser
		755					760					765			
Ala		Arg	Leu	Gln	Ala		Glu	Ala	Ala	His		Ala	Ala	Gly	Pro
	770					775	_				780		_		
	Glu	Val	Leu	Ala		Glu	Pro	Pro	Ala		Arg	Ala	Pro	Arg	
785					790					795					800
															<i>m</i>
Thr	Arg	Ala	Pro		Ala	Ser	Pro	Ala		Leu	Gly	Ser	Thr	Ala	Thr
	6	D		805	D				810	01		61		815	61
Ala	Ser	Pro		Ala	Pro	Ala	Arg		Arg	Gly	Leu	Glu		Ser	Glu
C	*		820	וכו	C	TI.	C1	825	C .	,	C1	V 1	830	Α.	Α.
Cys	Lys		Leu	rne	Ser	Inr		ınr	5er	Leu	GIn		HIS	Arg	Arg
11.	11.2 -	835	C1	C1	Λ	D	840	D	C	D	A	845	C1	1	A 1
116		Inr	Gly	610	Arg	855	1 V I	Pro	Cys	rro		Cys	61 y	Lys	V19
Dlac	850	Cl.	Com	The	u; a		1	A = 10	u; a	A 24.2	860	Lau	D; a	The	C1v
	Arg	GIII	ser	1111		rea	LyS	ASP	nis		Arg	Leu	ms	Thr	
865	Ara	Dro	Dha	ΛΙο	870 Cvs	61	Vol	Cva	C1v	875	A 1 c	Dha	A16	He	880
01 u	ui g	110	1116	885	Cys	0.10	101	CyS	890	LyS	VIS	ine	ліа	895	JEI
Mot	Ara	Lov	Ala		Hic	Ara	Arra	110		Tha	Clv	Clu	Δεσ	Pro	Tur
we r	ur g	Leu	mid	01 u	11113	ur 8	mg	116	1112	1111	OIA	OIU	ui g	110	1 7 1

900 905 910 Ser Cys Pro Asp Cys Gly Lys Ser Tyr Arg Ser Phe Ser Asn Leu Trp 925 920 Lys His Arg Lys Thr His Gln Gln His Gln Ala Ala Val Arg Gln 930 935 940 Gln Leu Ala Glu Ala Glu Ala Ala Val Gly Leu Ala Val Met Glu Thr 950 955 Ala Val Glu Ala Leu Pro Leu Val Glu Ala Ile Glu Ile Tyr Pro Leu 965 970 975 Ala Glu Ala Glu Gly Val Gln Ile Ser Gly 980 985

<210> 4700

<211> 441

<212> PRT

<213> Homo sapiens

<400> 4700

Met Glu Tyr Asp Glu Lys Leu Ala Arg Phe Arg Gln Ala His Leu Asn

1 5 10 15

Pro Phe Asn Lys Gln Ser Gly Pro Arg Gln His Glu Gln Gly Pro Gly

20 25 30

Glu Glu Val Pro Asp Val Thr Pro Glu Glu Ala Leu Pro Glu Leu Pro
35 40 45

Pro Gly Glu Pro Glu Phe Arg Cys Pro Glu Arg Val Met Asp Leu Gly 50 55 60

Leu Ser Glu Asp His Phe Ser Arg Pro Val Leu Arg Gln Ala Ile Glu 65 70 75 80

Glu Cys Lys Gln Val 11e Leu Glu Leu Pro Glu Gln Ser Glu Lys Gln 85 90 95

Lys Asp Ala Val Val Arg Leu Ile His Leu Arg Leu Lys Leu Gln Glu 100 105 110

Leu Lys Asp Pro Asn Glu Asp Glu Pro Asn Ile Arg Val Leu Leu Glu 115 120 125

His Arg Phe Tyr Lys Glu Lys Ser Lys Ser Val Lys Gln Thr Cys Asp

	130					135					140				
Lys	Cys	Asn	Thr	Ile	lle	Trp	G1 y	Leu	He	Gln	Thr	Trp	Tyr	Thr	Cys
145					150					155					160
Thr	G]y	Cys	Tyr	Tyr	Arg	Cys	His	Ser	Lys	Cys	Leu	Asn	Leu	Ile	Ser
-				165					170					175	
Lys	Pro	Cys	Val	Ser	Ser	Lys	Val	Ser	His	Gln	Ala	Glu	Tyr	Glu	Leu
			180					185					190		
Asn	lle	Cys	Pro	Glu	Thr	Gly	Leu	Asp	Ser	Gln	Asp	Tyr	Arg	Cys	Ala
		195					200					205			
Glu	Cys	Arg	Val	Pro	Ile	Ser	Leu	Arg	Gly	Val	Pro	Şer	Glu	Ala	Arg
	210					215					220				
Gln	Cys	Asp	Tyr	Thr	Gly	Gln	Tyr	Tyr	Cys	Ser	His	Cys	His	Trp	Asn
225					230					235					240
Asp	Leu	Ala	Val	He	Pro	Ala	Arg	Val	Val	His	Asn	Trp	Asp	Phe	Glu
				245					250					255	
Pro	Arg	Lys	Val	Ser	Arg	Cys	Ser	Met	Arg	Tyr	Leu	Ala	Leu	Met	Val
			260					265					270		
Ser	Arg	Pro	Val	Leu	Arg	Leu	Arg	Glu	lle	Asn	Pro	Leu	Leu	Phe	Ser
		275					280					285			
Tyr	Val	Glu	Glu	Leu	Val	Glu	He	Arg	Lys	Leu	Arg	Gln	Asp	He	Leu
	290					295					300				
Leu	Met	Lys	Pro	Tyr	Phe	He	Thr	Cys	Arg	Glu	Ala	Met	Glu	Ala	Arg
305					310					315					320
Leu	Leu	Leu	Gln	Leu	Gln	Asp	Arg	Gln	His	Phe	Val	Glu	Asn	Asp	Glu
				325					330					335	
Met	Tyr	Ser	Val	Gln	Asp	Leu	Leu	Asp	Val	His	Ala	Gly	Arg	Leu	Gly
			340					345					350		
Cys	Ser		Thr	Glu	lle	His	Thr	Leu	Phe	Ala	Lys		lle	Lys	Leu
		355					360					365			
Asp		Glu	Λrg	Cys	Gln		Lys	Gly	Phe	Va]		Glu	Leu	Cys	Arg
	370					375					380				
	Gly	Asp	Val	Leu		Pro	Phe	Asp	Ser		Thr	Ser	Va1	Cys	
385					390					395					400
Asp	Cys	Ser	Ala		Phe	His	Arg	Asp		Tyr	Tyr	Asp	Asn		Thr
				405					410					415	
Thr	Cys	Pro	Lys	Cys	Ala	Arg	Leu	Ser	Leu	Arg	Lys	Gln	Ser	Leu	Phe

Gln Glu Pro Gly Pro Asp Val Glu Ala <210> 4701 <211> 1389 <212> PRT <213> Homo sapiens <400> 4701 Met Lys Gln Leu Gln Pro Gln Pro Pro Pro Lys Met Gly Asp Phe Tyr Asp Pro Glu His Pro Thr Pro Glu Glu Glu Glu Asn Glu Ala Lys Ile Glu Asn Val Gln Lys Thr Gly Phe Ile Lys Gly Pro Met Phe Lys Gly Val Ala Ser Ser Arg Phe Leu Pro Lys Gly Thr Lys Thr Lys Val Asn Leu Glu Glu Gln Gly Arg Gln Lys Val Ser Phe Ser Phe Ser Leu Thr Lys Lys Thr Leu Gln Asn Arg Phe Leu Thr Ala Leu Gly Asn Glu Lys Glń Ser Asp Thr Pro Asn Pro Pro Ala Val Pro Leu Gln Val Asp Ser Thr Pro Lys Met Lys Met Glu Ile Gly Asp Thr Leu Ser Thr Ala Glu Glu Ser Ser Pro Pro Lys Ser Arg Val Glu Leu Gly Lys lle His Phe Lys Lys His Leu Leu His Val Thr Ser Arg Pro Leu Leu Ala Thr Thr Thr Ala Val Ala Ser Pro Pro Thr His Ala Ala Pro Leu Pro Ala Val

lle Ala Glu Ser Thr Thr Val Asp Ser Pro Pro Ser Ser Pro Pro Pro

Pro Pro Pro Pro Ala Gln Ala Thr Thr Leu Ser Ser Pro Ala Pro Val

		195				•	200					205			
Thr	Glu	Pro	Val	Ala	Leu	Pro	His	Thr	Pro	He	Thr	Val	Leu	Met	Ala
	210					215					220				
Ala	Pro	Val	Pro	Leu	Pro	Va]	Asp	Val	Ala	Val	Arg	Ser	Leu	Lys	Glu
225					230					235					240
Pro	Pro	He	He	He	Val	Pro	Glu	Ser	Leu	Glu	Ala	Asp	Thr	Lys	Gln
				245					250					255	
Asp	Thr	lle	Ser	Asn	Ser	Leu	Glu	Glu	His	Val	Thr	Gln	Ile	Leu	Asn
			260					265					270		
Glu	Gln	Ala	Asp	Ile	Ser	Ser	Lys	Lys	Glu	Asp	Ser	His	Ile	Gly	Lys
		275					280					285			
Asp	Glu	Glu	lle	Pro	Asp	Ser	Ser	Lys	lle	Ser	Leu	Ser	Cys	Lys	Lys
	290					295					300				
Thr	Gly	Ser	Lys	Lys	Lys	Ser	Ser	Gln	Ser		G1 y	He	Phe	Leu	
305					310					315					320
Ser	Glu	Ser	Asp		Asp	Ser	Va]	Arg		Ser	Ser	Ser	Gln		Ser
				325					330					335	
His	Asp	Leu		Phe	Ser	Ala	Ser		Glu	Lys	Glu	Arg	Asp	Phe	Lys
			340		_		_	345					350		
Lys	Ser		Ala	Pro	Leu	Lys		Glu	Asp	Leu	GIy		Pro	Ser	Arg
C		355					360	T	DI	C	т	365	,	,	C1
Ser		Inr	Asp	Arg	Asp		Lys	lyr	Pne	Ser			Lys	Leu	GIU
Δ	370	ТЬ	Λ	Т	V.a.1	375	Car	A ~	Cua	A 20 cm	380	•	A 22.00	C1	A 111 cm
	ASP	ınr	Arg	LÀL	390	261.	ser	Arg	Cys	395	Set.	Glu	Arg	Glu	400
385	Δra	Sor	Ara	Sor		Sor	Ara	Sor	Glu		Glv	Sor	Arg	Thr	
Mg	Mg	361	Mg	405		361	m g		410					415	
Len	Ser	Tvr	Ser			Glu	Arø						Ser		
БСС	501	1,1	420	8	001	0.4	8	425		.,.	.,1	пор	430	пор	6
Arg	Tvr	His		Ser	Ser	Pro	Tyr		Glu	Arg	Thr	Arg	Tyr	Ser	Gln
6	- , -	435	6				440	0		0		445	- , -		
Pro	Tyr	Thr	Asp	Asn	Arg	Ala	Arg	G]u	Ser	Ser	Asp	Ser	Glu	Glu	Glu
	450					455					460				
Tyr		Lys	Thr	Tyr	Ser	Arg	Arg	Thr	Ser	Ser	His	Ser	Ser	Ser	Tyr
465					470					475					480
Λrσ	Asp	Len	Arg	Thr	Ser	Ser	Tyr	Ser	Lvs	Ser	Asp	Arg	Asp	Cvs	Lvs

				485					490					495	
Thr	Glu	Thr	Ser	Tyr	Leu	Glu	Met	Glu	Arg	Arg	Gly	Lys	Tyr	Ser	Ser
			500					505					510		
Lys	Leu	Glu	Arg	G1u	Ser	Lys	Arg	Thr	Ser	Glu	Asn	G1u	Ala	Ile	Lys
		515					520					525			
Arg	Cys	Cys	Ser	Pro	Pro	Asn	Glu	Leu	Gly	Phe	Arg	Arg	G1 y	Ser	Ser
	530					535					540				
Tyr	Ser	Lys	His	Asp	Ser	Ser	Ala	Ser	Arg	Tyr	Lys	Ser	Thr	Leu	Ser
545					550					555					560
Lys	Pro	Ile	Pro	Lys	Ser	Asp	Lys	Phe	Lys	Asn	Ser	Phe	Cys	Cys	Thr
				565					570					575	
Glu	Leu	Asn	Glu	Glu	lle	Lys	Gln	Ser	His	Ser	Phe	Ser	Leu	Gln	Thr
			580					585					590		
Pro	Cys	Ser	Lys	Gly	Ser	Glu	Leu	Arg	Met	He	Asn	Lys	Asn	Pro	Glu
		595					600					605			
Arg	Glu	Lys	Ala	61 y	Ser	Pro	Ala	Pro	Ser	Asn	Arg	Leu	Asn	Asp	Ser
	610					615					620				
Pro	Thr	Leu	Lys	Lys	Leu	Asp	Glu	Leu	Pro	Ile	Phe	Lys	Ser	Glu	Phe
625					630					635					640
Ile	Thr	His	Asp	Ser	His	Asp	Ser	Ile	Lys	Glu	Leu	Asp	Ser	Leu	Ser
				645					650					655	
Lys	Val	Lys	Asn	Asp	Gln	Leu	Arg	Ser	Phe	Cys	Pro	lle	Glu	Leu	Asn
			660					665					670		
lle	Asn	Gly	Ser	Pro	Gly	Ala	Glu	Ser	Asp	Leu	Ala	Thr	Phe	Cys	Thr
		675					680					685			
Ser	Lys	Thr	Asp	Ala	Val	Leu	Met	Thr	Ser	Asp	Asp	Ser	Val	Thr	Gly
	690					695					700				
Ser	Glu	Leu	Ser	Pro	Leu	Val	Lys	Ala	Cys	Met	Leu	Ser	Ser	Asn	Gly
705					710					715					720
Phe	Gln	Asn	He	Ser	Arg	Cys	Lys	Glu	Lys	Лѕр	Leu	Asp	Λsp	Thr	Cys
				725					730					735	
Met	Leu	His	Lys	Lys	Ser	Glu	Ser	Pro	Phe	Arg	Glu	Thr	Glu	Pro	Leu
			740					745					750		
Val	Ser	Pro	His	Gln	Asp	Lys	Leu	Met	Ser	Met	Pro	Val	Met	Thr	Val
		755					760					765			
Asp	Tyr	Ser	Lys	Thr	Val	Val	Lys	Glu	Pro	Val	Asp	Thr	Arg	Val	Ser

	770					775					780				
Cys	Cys	Lys	Thr	Lys	Asp	Ser	Asp	He	Tyr	Cys	Thr	Leu	Asn	Asp	Ser
785					790					795					800
Asn	Pro	Ser	Leu	Cys	Asn	Ser	Glu	Ala	Glu	Asn	He	Glu	Pro	Ser	Val
				805					810					815	
Met	Lys	Пе	Ser	Ser	Asn	Ser	Phe	Met	Asn	Val	His	Leu	Glu	Ser	Lys
			820					825					830		
Pro	Va]	He	Cys	Asp	Ser	Arg	Asn	Leu	Thr	Asp	His	Ser	Lys	Phe	Ala
		835					840					845			
Cys	Glu	Glu	Tyr	Lys	Gln	Ser	Ile	Gly	Ser	Thr	Ser	Ser	Ala	Ser	Val
	850					855					860				
Asn	His	Phe	Asp	Asp	Leu	Tyr	Gln	Pro	lle	Gly	Ser	Ser	Gly	lle	Ala
865					870					875					880
Ser	Ser	Leu	Gln	Ser	Leu	Pro	Pro	Gly	He	Lys	Val	Asp	Ser	Leu	Thr
				885					890					895	
Leu	Leu	Lys	Cys	Gly	Glu	Asn	Thr	Ser	Pro	Val	Leu	Asp	Ala	Val	Leu
			900					905					910		
Lys	Ser	Lys	Lys	Ser	Ser	Glu	Phe	Leu	·Lys	His	Ala	G1 y	Lys	Glu	Thr
		915					920					925			
He	Val	Glu	Val	Gly	Ser	Asp	Leu	Pro	Asp	Ser	Gly	Lys	Gly	Phe	Ala
	930					935					940				
Ser	Arg	Glu	Asn	Arg	Arg	Asn	Asn	Gly	Leu	Ser	Gly	Lys	Cys	Leu	Gln
945					950					955					960
Glu	Ala	G1n	Glu	Glu	Gly	Asn	Ser	lle	Leu	Pro	Glu	Arg	Arg	Gly	Arg
				965					970					975	
Pro	Glu	He	Ser	Leu	Asp	Glu	Arg	Gly	Glu	G]y	Gly	His	Val	His	Thr
			980					985					990		
Ser	Asp	Asp	Ser	Glu	Val	Val	Phe	Ser	Ser	Cys			Asn	Leu	Thr
		995					1000					1005			
		Asp	Ser	Asp			Thr	Tyr	Λla			Cys	Asp	Ser	Ser
	1010					1015	_				1020	0.0	_	_	
		Ala	Pro			Val	Ser	Thr			Glu	Asp	Tyr		Gly
102					1030			_		1035	0.1		mı.		1040
Ser	Ser	G]u			Asn	Asp	Glu			Ser	Glu	Asp			Ser
		c		1045	D				1050	61	C	17 7		1055	17 1
Asp	Asp	Ser	5er	He	Pro	Arg	Asn	Arg	Leu	GIn	Ser	val	val	val	val

	1060			1	065			1	070		
Pro Lys As	n Ser	Thr Le	eu Pro	Met	Glu Gl	u Thr	Ser	Pro	Cys	Ser	Ser
107	5		1	080			J	085			
Arg Ser Se	r Gln	Ser T	yr Arg	His	Tyr Se	r Asp	His	Trp	Glu.	Asp	Glu
1090			1095]	100				
Arg Leu Gl	u Ser	Arg A	rg His	Leu	Tyr Gl	u Glu	Lys	Phe	Glu	Ser	11e
1105		11	10			1115				1	120
Ala Ser Ly	s Ala	Cys P	ro Gln	Thr	Asp Ly	s Phe	Phe	Leu	His	Lys	Gly
		1125			113	0			1	135	
Thr Glu Ly	s Asn	Pro G	lu Ile	Ser	Phe Th	r Gln	Ser	Ser	Arg	Lys	Gln
	1140			1	145			1	1150		
lle Asp As	n Arg	Leu P	ro Glu	Leu	Ser Hi	s Pro	Gln	Ser	Asp	Gly	Val
115	55		1	160			1	165			
Asp Ser Th	ır Ser	His T	hr Asp	Val	Lys Se	r Asp	Pro	Leu	Gly	His	Pro
1170			1175				1180				
Asn Ser G	u Glu	Thr V	al Lys	Ala	Lys Il	e Pro	Ser	Arg	Gln	G1n	Glu
1185		11	90			1195		•		l	200
Glu Leu Pr	o Ile	Tvr S	er Ser	Asp	Phe G1	u Asp	Phe	Pro	Asn	Lys	Ser
		. ,	0. 00.			p				-	
		1205	0. 00.		121					215	
Trp Gln G		1205			121	0			1	215	
		1205		Asn	121	0		Arg	1	215	
	n Thr 1220	1205 Thr P	he Gln	Asn 1	121 Arg Pr 225	0 o Asp	Ser	Arg	1 Leu 1230	215 Gly	Lys
Trp Gln G	n Thr 1220 eu Ser	1205 Thr P	he Gln er Ser	Asn 1	121 Arg Pr 225	0 o Asp	Ser Pro	Arg	1 Leu 1230	215 Gly	Lys
Trp Gln G.	n Thr 1220 eu Ser 35	1205 Thr P Phe S	he Gln er Ser	Asn I Ser 1240	121 Arg Pr 225 Cys Gl	o Asp	Ser Pro	Arg His 1245	1 Leu 1230 Val	215 Gly Asp	Lys Gly
Trp Gln G. Thr Glu Lo	n Thr 1220 eu Ser 35	1205 Thr P Phe S	he Gln er Ser	Asn I Ser 1240	121 Arg Pr 225 Cys Gl	o Asp u lle u Gly	Ser Pro	Arg His 1245	1 Leu 1230 Val	215 Gly Asp	Lys Gly
Trp Gln G. Thr Glu Le 123 Leu His Se	n Thr 1220 eu Ser 35 er Ser	Thr P Phe S Glu G	he Gln er Ser Hu Leu 1255	Asn l Ser 1240 Arg	121 Arg Pr 225 Cys G1 Asn Le	o Asp u lle u Gly	Ser Pro Trp	Arg His 1245 Asp	Leu 1230 Val Phe	215 Gly Asp Ser	Lys Gly Gln
Trp Gln G. Thr Glu Le 123 Leu His Se 1250	n Thr 1220 eu Ser 35 er Ser	Thr P Phe S Glu G Thr T	he Gln er Ser Hu Leu 1255	Asn 1 Ser 1240 Arg Gln	121 Arg Pr 225 Cys G1 Asn Le	o Asp u lle u Gly	Ser Pro Trp 1260 Ser	Arg His 1245 Asp Ser	Leu 1230 Val Phe	215 Gly Asp Ser	Lys Gly Gln
Trp Gln G. Thr Glu Le 12: Leu His Se 1250 Glu Lys Pr	n Thr 1220 eu Ser 35 er Ser	1205 Thr P Phe S Glu G Thr T 12	he Gln er Ser flu Leu 1255 hr Tyr	Asn l Ser 1240 Arg Gln	121 Arg Pr 225 Cys Gl Asn Le	o Asp u lle u Gly o Asp 1275	Pro Trp 1260 Ser	Arg His 1245 Asp Ser	Leu 1230 Val Phe Tyr	215 Gly Asp Ser Gly	Lys Gly Gln Ala 280
Trp Gln G. Thr Glu Le 123 Leu His Se 1250 Glu Lys Pr 1265	n Thr 1220 eu Ser 85 er Ser co Ser	1205 Thr P Phe S Glu G Thr T 12	he Gln er Ser flu Leu 1255 hr Tyr	Asn l Ser 1240 Arg Gln	121 Arg Pr 225 Cys Gl Asn Le	o Asp u lle u Gly o Asp 1275 a Glu	Pro Trp 1260 Ser	Arg His 1245 Asp Ser	Leu 1230 Val Phe Tyr	215 Gly Asp Ser Gly	Lys Gly Gln Ala 280
Trp Gln G. Thr Glu Le 123 Leu His Se 1250 Glu Lys Pr 1265	n Thr 1220 eu Ser 85 er Ser co Ser	1205 Thr P Phe S Glu G Thr T 12 Lys T	he Gln Fer Ser Flu Leu 1255 Fhr Tyr F70 Fyr Gln	Asn l Ser 1240 Arg Gln	121 Arg Pr 225 Cys G1 Asn Le G1n Pr Asn A1	o Asp u lle u Gly o Asp 1275 a Glu	Ser Pro Trp 1260 Ser Gln	Arg His 1245 Asp Ser	Leu 1230 Val Phe Tyr Gly	215 Gly Asp Ser Gly 1 Gly 295	Lys Gly Gln Ala 280 Thr
Trp Gln G. Thr Glu Le 123 Leu His Se 1250 Glu Lys Pr 1265 Cys Gly G	n Thr 1220 eu Ser 85 er Ser co Ser	1205 Thr P Phe S Glu G Thr T 12 Lys T 1285 Gln G	he Gln Fer Ser Flu Leu 1255 Fhr Tyr F70 Fyr Gln	Asn I Ser I240 Arg Gln Gln	121 Arg Pr 225 Cys G1 Asn Le G1n Pr Asn A1	o Asp u lle u Gly o Asp 1275 a Glu	Ser Pro Trp 1260 Ser Gln	Arg His 1245 Asp Ser Tyr	Leu 1230 Val Phe Tyr Gly	215 Gly Asp Ser Gly 1 Gly 295	Lys Gly Gln Ala 280 Thr
Trp Gln G. Thr Glu Le 123 Leu His Se 1250 Glu Lys Pr 1265 Cys Gly G	n Thr 1220 eu Ser 35 er Ser o Ser y His	Thr P. Phe S Glu G Thr T 12 Lys T 1285 Gln G	the Gln for Ser flu Leu flu Leu flu Leu flu Tyr fru fru fru flu Gln fly Asn	Asn I Ser 1240 Arg Gln Gln Gly	Arg Pr 225 Cys Gl Asn Le Gln Pr Asn Al 129 Tyr Tr 305	o Asp u lle u Gly o Asp 1275 a Glu o	Ser Pro Trp 1260 Ser Gln Pro	Arg His 1245 Asp Ser Tyr	Leu 1230 Val Phe Tyr Gly 1 Ser	215 Gly Asp Ser Gly 1 Gly 295 Gly	Lys Gly Gln Ala 280 Thr
Trp Gln G. Thr Glu Le 12: Leu His Se 1250 Glu Lys Pr 1265 Cys Gly G	n Thr 1220 Pu Ser S5 Pr Ser Vr Trp 1300 ly Thr	Thr P. Phe S Glu G Thr T 12 Lys T 1285 Gln G	the Gln fer Ser flu Leu 1255 fhr Tyr f70 fyr Gln fily Asn	Asn I Ser 1240 Arg Gln Gln Gly	Arg Pr 225 Cys Gl Asn Le Gln Pr Asn Al 129 Tyr Tr 305	o Asp u lle u Gly o Asp 1275 a Glu o	Ser Pro Trp 1260 Ser Gln Pro	Arg His 1245 Asp Ser Tyr	Leu 1230 Val Phe Tyr Gly 1 Ser	215 Gly Asp Ser Gly 1 Gly 295 Gly	Lys Gly Gln Ala 280 Thr
Trp Gln G. Thr Glu Le 12: Leu His Sc 1250 Glu Lys Pr 1265 Cys Gly G Arg Asp Ty	n Thr 1220 Pu Ser 35 Pr Ser To Ser 1300 1y Thr	Thr P Phe S Glu G Thr T 12 Lys T 1285 Gln G Gly V	the Gln er Ser flu Leu 1255 Thr Tyr 70 fyr Gln fly Asn /al Val	Asn I Ser 1240 Arg Gln Gln Gly I Tyr 1320	Arg Pr 225 Cys Gl Asn Le Gln Pr Asn Al 129 Tyr Tr 305 Asp Ar	o Asp u lle u Gly to Asp 1275 a Glu o rp Asp	Ser Pro Trp 1260 Ser Gln Pro	Arg His 1245 Asp Ser Tyr Arg Gly 1325	Leu 1230 Val Phe Tyr Gly 1 Ser 1310 Gln	215 Gly Asp Ser Gly 295 Gly Val	Lys Gly Gln Ala 280 Thr Arg
Trp Gln G. Thr Glu Le 123 Leu His Se 1250 Glu Lys Pr 1265 Cys Gly G. Arg Asp Ty Pro Pro G.	n Thr 1220 Pu Ser 35 Pr Ser To Ser 1300 1y Thr	Thr P Phe S Glu G Thr T 12 Lys T 1285 Gln G Gly V	the Gln er Ser flu Leu 1255 Thr Tyr 70 fyr Gln fly Asn /al Val	Asn I Ser 1240 Arg Gln Gln Gly I Tyr 1320	Arg Pr 225 Cys Gl Asn Le Gln Pr Asn Al 129 Tyr Tr 305 Asp Ar	o Asp u Ile u Gly o Asp 1275 a Glu o p Asp	Ser Pro Trp 1260 Ser Gln Pro	Arg His 1245 Asp Ser Tyr Arg Gly 1325	Leu 1230 Val Phe Tyr Gly 1 Ser 1310 Gln	215 Gly Asp Ser Gly 295 Gly Val	Lys Gly Gln Ala 280 Thr Arg

1345 1350 1355 1360

Gln Lys Asp Lys Gly Ser Val Gln Ala Pro Glu Ile Ser Ser Asn Ser

1365 1370 1375

Ile Lys Asp Thr Leu Ala Val Asn Glu Lys Lys Asp Phe

1380 1385

<210> 4702

<211> 134

<212> PRT

<213> Homo sapiens

<400> 4702

Met Val Gly Leu Leu Phe His Ala Pro Lys Ala Pro Glu Met Ala Pro
1 5 10 15

Leu Arg Cys Cys lle Met Asn Lys Ile Ile Met Val Arg Arg Pro Lys 20 25 30

Gln Ser Thr Ala Asp Tyr Gly Met Arg Thr Ser Gly Pro Val Glu Ser 35 40 45

Gly Leu Ser Ala Asp Ser Leu Gln Leu Leu Cys Ser Tyr Ala Ala Ile 50 55 60

Lys Asn Ser Ala Glu Leu Leu Met Val Gly Pro Gln Gly Met Arg Pro 65 70 75 80

Ala Thr Gly Gln Asp Leu Leu Cys Arg Pro Cys Leu Ser His Asp Pro
85 90 95

Pro Gly Pro Leu His Pro Pro Arg Gly Leu Ser Gly Ser Ser Ser Leu 100 105 110

Leu Ile Ser Pro Arg Leu Gln Asp Val Ser Leu Gln Leu Val His Pro 115 120 125

Thr Pro Glu Glu Ser Phe

130

<210> 4703

<211> 743

<212	2> PF	RT													
<213	3> Hc	omo s	sapie	ens											
<400)> 47	703													
Met	Ser	Ser	Leu	Ala	Ala	Lys	Arg	Leu	Gly	Met	Asn	Arg	Arg	Pro	Ala
1				5					10					15	
Gly	Ser	Gly	Gly	G1 y	Gly	Gly	Glu	Ala	Ala	Thr	Trp	Gly	His	Arg	Phe
			20					25					30		
Trp	Arg	Pro	Gln	Glu	Arg	Pro	Thr	Asp	Arg	Asn	Gln	G1 y	Glu	Met	Ala
		35					40					45			
His	Thr	Cys	Arg	Gly	Thr	Ile	Asn	Leu	Ser	Thr	Ala	His	Пе	Asp	Thr
	50					55					60				
Glu	Asp	Ser	Cys	Gly	He	Leu	Leu	Thr	Ser	Gly	Ala	Arg	Ser	Tyr	His
65					70					75					80
Leu	Lys	Ala	Ser	Ser	Glu	Val	Asp	Arg	Gln	Gln	Trp	Ile	Thr	Ala	Leu
				85					90					95	
Glu	Leu	Ala	Lys	Ala	Lys	Ala	Val	Arg	Val	Met	Asn	Thr	His	Ser	Asp
			100					105					110		
Asp	Ser	Gly	Asp	Asp	Asp	Glu	Ala	Thr	Thr	Pro	Ala	Asp	Lys	Ser	Glu
		115					120					125			
Leu	His	His	Thr	Leu	Lys	Asn	Leu	Ser	Leu	Lys	Leu	Asp	Asp	Leu	Ser
	130					135					140				
Thr	Cys	Asn	Asp	Leu	He	Ala	Lys	His	Gly	Ala	Ala	Leu	Gln	Arg	Ser
145					150					155					160
Leu	Thr	Glu	Leu	Asp	Gly	Leu	Lys	11e	Pro	Ser	Glu	Ser	G1 y	Glu	Lys
				165					170					175	•
Leu	Lys	Val	Val	Asn	Glu	Arg	Ala	Thr	Leu	Phe	Arg	He	Thr	Ser	Asn
			180					185					190		
Ala	Met	He	Asn	Ala	Cys	Arg	Asp	Phe	Leu	Glu	Leu	Ala	Glu	lle	His
		195					200					205			
Ser	Arg	Lys	Trp	Gln	Arg	Ala	Leu	Gln	Tyr	Glu	Gln	Glu	Gln	Arg	Val
	210					215					220				
His	Leu	G1u	Glu	Thr	Пe	Glu	Gln	Leu	Ala	Lys	Gln	His	Asn	Ser	Leu

Glu Arg Ala Phe His Ser Ala Pro Gly Arg Pro Ala Asn Pro Ser Lys

Ser	Phe	lle	Glu	Gly	Ser	Leu	Leu	Thr	Pro	Lys	Gly	Glu	Asp	Ser	Glu
			260					265					270		
Glu	Asp	Glu	Asp	Thr	Glu	Tyr	Phe	Asp	Ala	Met	Glu	Asp	Ser	Thr	Ser
		275					280					285			
Phe	He	Thr	Val	lle	Thr	Glu	Ala	Lys	Glu	Asp	Ser	Arg	Lys	Ala	Glu
	290					295					300				
Gly	Ser	Thr	Gly	Thr	Ser	Ser	Val	Asp	Trp	Ser	Ser	Ala	Asp	Asn	Val
305					310					315					320
Leu	Asp	Gly	Ala	Ser	Leu	Val	Pro	Lys	Gly	Ser	Ser	Lys	Val	Lys	Arg
				325					330					335	
Arg	Val	Arg	lle	Pro	Asn	Lys	Pro	Asn	Tyr	Ser	Leu	Asn	Leu	Trp	Ser
			340					345					350		
lle	Met	Lys	Asn	Cys	lle	Gly	Arg	Glu	Leu	Ser	Arg	He	Pro	Met	Pro
		355					360					365			
Val	Asn	Phe	Asn	Glu	Pro	Leu	Ser	Met	Leu	Gln	Arg	Leu	Thr	Glu	Asp
	370					375					380				
Leu	Glu	Tyr	His	His	Leu	Leu	Asp	Lys	Ala	Val	His	Cys	Thr	Ser	Ser
385					390					395					400
Val	Glu	Gln	Met	Cys	Leu	Va]	Ala	Ala	Phe	Ser	Val	Ser	Ser	Tyr	Ser
				405					410					415	
Thr	Thr	Val	His	Arg	lle	Ala	Lys	Pro	Phe	Asn	Pro	Met	Leu	Gly	Glu
			420					425					430		
Thr	Phe	Glu	Leu	Asp	Arg	Leu	Asp	Asp	Met	Gly	Leu	Arg	Ser	Leu	Cys
		435					440					445			
Glu	Gln	Val	Ser	His	His	Pro	Pro	Ser	Ala	Ala	His	Tyr	Val	Phe	Ser
	450					455					460				
Lys	His	Gly	Trp	Ser	Leu	Trp	Gln	Glu	He	Thr	He	Ser	Ser	Lys	Phe
465					470					475					480
Arg	Gly	Lys	Tyr	11e	Ser	He	Met	Pro	Leu	Gly	Ala	He	His	Leu	Glu
				485					490					495	
Phe	Gln	Ala	Ser	G1 y	Asn	His	Tyr	Val	Trp	Arg	Lys	Ser	Thr	Ser	Thr
			500					505					510		
Val	His	Asn	lle	He	Val	61y	Lys	Leu	Trp	lle	Asp	Gln	Ser	Gly	Asp
		515					520					525			
lle	Glu	He	Val	Asn	His	Lys	Thr	Asn	Asp	Arg	Cys	Gln	Leu	Lys	Phe
	530					535					540				

Leu Pro Tyr Ser Tyr Phe Ser Lys Glu Ala Ala Arg Lys Val Thr Gly 550 555 545 Val Val Ser Asp Ser Gln Gly Lys Ala His Tyr Val Leu Ser Gly Ser 570 575 565 Trp Asp Glu Gln Met Glu Cys Ser Lys Val Met His Ser Ser Pro Ser 580 585 Ser Pro Ser Ser Asp Gly Lys Gln Lys Thr Val Tyr Gln Thr Leu Ser 600 Ala Lys Leu Leu Trp Lys Lys Tyr Pro Leu Pro Glu Asn Ala Glu Asn 615 610 Met Tyr Tyr Phe Ser Glu Leu Ala Leu Thr Leu Asn Glu His Glu Glu 630 635 Gly Val Ala Pro Thr Asp Ser Arg Leu Arg Pro Asp Gln Arg Leu Met 650 655 645 Glu Lys Gly Arg Trp Asp Glu Ala Asn Thr Glu Lys Gln Arg Leu Glu 670 660 665 Glu Lys Gln Arg Leu Ser Arg Arg Arg Arg Leu Glu Ala Cys Gly Pro . 680 685 Gly Ser Ser Cys Ser Ser Glu Glu Glu Lys Glu Ala Asp Ala Tyr Thr 695 690 Pro Leu Trp Phe Glu Lys Arg Leu Asp Pro Leu Thr Gly Glu Met Ala 710 715 Cys Val Tyr Lys Gly Gly Tyr Trp Glu Ala Lys Glu Lys Gln Asp Trp 725 735 730 His Met Cys Pro Asn Ile Phe 740

<210> 4704

<211> 577

<212> PRT

<213> Homo sapiens

<400> 4704

Met Glu Asn Glu Arg Thr Lys Asp Leu lle lle Glu Gln Arg Phe His

1 5 10 15

Arg	Thr	He	lle	Gly	Gln	Lys	Gly	Glu	Arg	lle	Arg	Glu	He	Arg	Asp
			20					25					30		
Lys	Phe	Pro	Glu	Val	lle	He	Asn	Phe	Pro	Asp	Pro	Ala	Gln	Lys	Ser
		35					40					45			
Asp	lle	Va]	Gln	Leu	Arg	Gly	Pro	Lys	Asn	Glu	Val	Glu	Lys	Cys	Thr
	50					55					60				
Lys	Tyr	Met	Gln	Lys	Met	Val	Ala	Asp	Leu	Val	Glu	Asn	Ser	Tyr	Ser
65					70					75					80
Ile	Ser	Val	Pro	He	Phe	Lys	Gln	Phe		Lys	Asn	lle	lle	G1 y	Lys
				85					90					95	
Gly	Gly	Ala		lle	Lys	Lys	Ile		Glu	Glu	Ser	Asn		Lys	He
		_	100					105					110		
Asp	Leu		Ala	Glu	Asn	Ser		Ser	Glu	Thr	He		He	Thr	Gly
		115			0.1		120					125	0	2.1	0.1
Lys		Ala	Asn	Cys	Glu		Ala	Arg	Ser	Arg		Leu	Ser	11e	Gin
,	130	1	4.1	A	T 1	135	C1	W - 1	C1	V. 1	140	11.	D	A 1 -	1
	Asp	Leu	Ala	Asn		Ala	GIU	vaı	610		ser	116	Pro	Ala	
145	uio	Aon	Cor	Lou	150	Clu	The	lva	Cly	155	1	Tla	Ana	Son	160
Leu	1115	ASII	261	165	116	GIY	1111	LyS	170	AI g	Leu	116	Alg	Ser 175	116
Mat	C1u	Glu	Cvs		G1v	Val	Hic	Πρ		Phe	Pro	Val	Glu	Gly	Ser
MCt	0.14	0,4	180	019	Oly	, 61	1113	185	11.15	1110	110	, 41	190	G.E.J	50,1
Glv	Ser	Asp		Val	Val	He	Arg		Pro	Ser	Ser	Asp		Glu	Lvs
V.,		195					200	**= 3				205			
Ala	Lvs		Gln	Leu	Leu	His		Ala	Glu	Glu	Lys		Thr	Lys	Ser
	210	•				215					220			•	
Phe		Val	Asp	lle	Arg	Ala	Lys	Pro	Glu	Tyr	His	Lys	Phe	Leu	He
225					230					235					240
Gly	Lys	Gly	Gly	Gly	Lys	lle	Arg	Lys	Val	Arg	Asp	Ser	Thr	G] y	Ala
				245					250					255	
Arg	Val	He	Phe	Pro	Ala	Ala	Glu	Asp	Lys	Asp	Gln	Asp	Leu	He	Thr
			260					265					270		
lle	He	Gly	Lys	G]u	Asp	Ala	Val	Arg	Glu	Ala	Gln	Lys	Glu	Leu	Glu
		275					280					285			
Ala	Leu	He	Gln	Asn	Leu	Asp	Asn	Val	Val	Glu	Asp	Ser	Met	Leu	Val
	290					295					300				

Asp	Pro	Lys	His	His	Arg	His	Phe	Val	Ile	Arg	Arg	Gly	Gln	Val	Leu
305					310					315					320
Arg	Glu	lle	Ala	Glu	Glu	Tyr	Gly	Gly	Val	Met	Val	Ser	Phe	Pro	Arg
				325					330					335	
Ser	Gly	Thr	Gln	Ser	Asp	Lys	Val	Thr	Leu	Lys	Gly	Ala	Lys	Asp	Cys
			340					345					350		
Val	Glu	Ala	Ala	Lys	Lys	۸rg	He	Gln	Glu	He	He	Glu	Asp	Leu	Glu
		355					360					365			
Ala	Gln	Val	Thr	Leu	Glu	Cys	Ala	lle	Pro	Gln	Lys	Phe	His	Arg	Ser
	370					375					380				
Val	Met	Gly	Pro	Lys	Gly	Ser	Arg	He	Gln	Gln	Ile	Thr	Arg	Asp	Phe
385					390					395					400
Ser	Val	Gln	He	Lys	Phe	Pro	Asp	Arg	Glu	Glu	Asn	Ala	Val	His	Ser
				405					410					415	
Thr	Glu	Pro	Val	Val	Gln	Glu	Asn	Gly	Asp	Glu	Ala	Gly	Glu	Gly	Arg
			420					425					430		
Glu	Ala	Lys	Asp	Cys	Asp	Pro	Gly	Ser	Pro	Arg	Arg	Cys	Asp	He	lle
		435					440					445			
Ile	Ile	Ser	Gly	Arg	Lys	Glu	Lys	Cys	Glu	Ala	Ala	Lys	Glu	Ala	Leu
	450					455					460				
Ġlu	Ala	Leu	Val	Pro	Val	Thr	Ile	Glu	Val	Glu	Val	Pro	Phe	Asn	Leu
465					470					475					480
His	Arg	Tyr	Val	lle	Gly	Gln	Lys	Gly	Ser	G1 y	He	Arg	Lys	Met	Met
				485					490					495	
Asp	Glu	Phe	Glu	Val	Asp	Pro	Phe	Pro	G1 y	Arg	Pro	Cys	His	Arg	Ser
			500					505					510		
Gly	Leu	Ser	His	Pro	Leu	Pro	Ser	Ala	Ser	Val	Leu	Ser	Gln	Leu	Pro
		515		-			520					525			
Val	Asp	Ser	Ala	Ser	Ser	Cys	Ser	Asp	Trp	Ala	Leu	Thr	Ser	Leu	His
	530					535					540				
G]y	Gly	Trp	Pro	Ser	Leu	Ser	Pro	Leu	Leu	Gly	Pro	Arg	Ala	Asn	Gly
545					550					555					560
Leu	Phe	Pro	Val	Leu	Trp	Gly	Pro	Glu	Arg	Gln	Va]	Ser	Pro	Leu	Phe
				565					570					575	

Thr

<211> 121 <212> PRT <213> Homo sapiens <400> 4705 Met Val Ala Asp Leu Ile Asn Val Gln Ile Thr Ser Gly Gln Ala Gln 10 1 Ser Ala Leu Phe Pro Pro Ser Pro Gln Leu Trp Leu Met Ser Ser Pro 25 Phe Leu Pro Trp Pro Arg His Gly Glu Ser Ser 11e Gly Lys Ala Lys 35 40 45 Val Ser Ser Arg Lys Thr Glu Gly Cys Ala Ser Pro Leu Glu Pro Leu 50 60 55 Pro Ser Trp Pro Gly Leu Leu Pro Val Ser Val Lys Lys Lys Lys 70 75 Lys Lys Lys Ser Asp Met Gln Leu Leu Ser Thr Cys Cys Met Ser Ser 85 90 Met Val Arg Asp Ala Glu Lys Tyr Lys Ala Val Ser Gly Gly Gln Val 105 110 Glu Tyr Gln Leu His His Leu Lys Thr 115 120 <210> 4706 <211> 202 <212> PRT <213> Homo sapiens

<210> 4705

<400> 4706

1

20 25 30

10

Met Arg Trp Gly Leu Val Pro Ser Trp Val Lys Glu Pro Lys Lys Phe

Thr Leu Leu lle Asn Ala Arg Ser Glu Thr Val Arg Asp Lys Pro Ala

Phe Lys Asn Ala Met Lys Arg Arg Val Leu Val Pro Ser Asp Gly Tyr Tyr Glu Trp Gln Asp Lys Asp Gly Arg Lys Arg Pro Phe Phe Ile 50 55 His Arg Arg Asp Gly Gln Pro Thr Gly Phe Ala Ala Leu Ala Glu Thr 70 75 Trp Met Gly Pro Asn Gly Glu Glu Phe Asp Ser Val Ala lle Val Thr 90 Thr Gln Ala Ser Pro Asp Leu Ala Glu Leu His His Arg Val Pro Val Thr Ile Ala Pro Asp Asp Phe Glu Arg Trp Leu Asp Gly Arg Ala Asn 120 Asp Val Glu Asp Val Met Pro Leu Leu Arg Ala Pro Arg Val Gly Glu 130 135 140 Phe Ala Trp His Glu Val Ser Thr Arg Val Asn Arg Val Ala Asn Asp 150 155 160 Asp Glu Gln Leu Val Leu Pro Ile Ser Glu Glu Gln Arg Ala Ala Glu 165 170 Ala Pro Lys Pro Val Lys Lys Ala Ala Pro Arg Lys Thr Thr Pro Glu 180 185 190 Pro Glu Asp Glu Gly Gln Gly Ser Leu Phe 195 200

<210> 4707

<211> 339

<212> PRT

<213> Homo sapiens

<400> 4707

 Met
 Leu
 Pro
 Ser
 Ala
 Val
 Ala
 Ala
 His
 Ala
 Gly
 Ala
 Tyr
 Trp
 Asp
 Val

 1
 5
 5
 10
 15
 15

 Val
 Ala
 Ser
 Ser
 Ala
 Leu
 Leu
 Asp
 Leu
 Pro
 Ala
 Ala
 Pro
 Gly
 Phe
 Gly

 Asn
 Leu
 Gly
 Lys
 Ser
 Phe
 Leu
 11e
 Glu
 Asn
 Leu
 Leu
 Arg
 Val
 Gly
 Gly

 35
 40
 45
 45

Ala	Pro	Thr	Pro	Arg	Leu	Gln	Pro	Pro	Ala	Pro	His	Asp	Pro	Ala	Thr
	50					55					60				
Ala	Leu	Ala	Thr	Ala	Gly	Ala	Gln	Leu	Arg	Pro	Leu	Pro	Ala	Ser	Pro
65					70					75					80
Va]	Pro	Leu	Lys	Leu	Cys	Pro	Ala	Ala	Glu	Gln	Val	Ser	Pro	Ala	Gly
				85					90					95	
Ala	Pro	Tyr	Gly	Thr	Arg	Trp	Ala	Phe	Gln	Val	Leu	Ser	Pro	Ser	Ala
			100					105					110		
Asp	Ser	Ala	Arg	Leu	Pro	Gly	Arg	Ala	Pro	Gly	Asp	Arg	Asp	Cys	Thr
		115					120					125			
Phe	Gln	Pro	Ser	Ala	Pro	Ala	Pro	Ser	Lys	Pro	Phe	Leu	Leu	Ser	Thr
	130					135					140				
Pro	Pro	Phe	Tyr	Ser	Ala	Cys	Cys	Gly	Gly	Ser	Cys	Arg	Arg	Pro	Ala
145					150					155					160
Ser	Ser	Thr	Ala	Phe	Pro	Arg	Glu	Glu	Ser	Val	Leu	Pro	Leu	Leu	Thr
				165					170					175	
Gln	Asp	Ser	Asn	Ser	Lys	Ala	Arg	Arg	Gly	Ile	Leu	Arg	Arg	Ala	Val
			180					185					190		
Phe	Ser	Glu	Asp	Gln	Arg	Lys	Ala	Leu	Glu	Lys	Met	Phe	Gln	Lys	Gln
		195					200					205			
Lys	Tyr	He	Ser	Lys	Thr	Asp	Arg	Lys	Lys	Leu	Ala	He	Asn	Leu	Gly
	210					215					220				
Leu	Lys	Glu	Ser	Gln	Val	Lys	lle	Trp	Phe	Gln	Asn	Arg	Arg	Met	Lys
225					230					235					240
Trp	Arg	Asn	Ser	Lys	Glu	Lys	Glu	Val	Leu	Ser	Asn	Arg	Cys	lle	Gln
				245					250					255	
Glu	Val	Gly		Gln	Glu	Asp	Pro		Ser	Arg	Ser	Ala		Gly	Phe
			260					265					270		
Pro	Ser		Cys	Pro	Ser	He		Asp	Val	Pro	Gln		His	Ser	Ser
		275					280					285			
Pro		Trp	Arg	Glu	Asn	Ser	Pro	Glu	Pro	Ser		Arg	Leu	He	Gln
	290				_	295					300				
	Ser	Ser	Gly	Ala		Pro	Pro	Glu	Ala		Ser	Leu	Gln	Gly	
305			_	_	310					315					320
Leu	Tyr	Leu	Cys	Ser	Glu	Glu	Glu	Ala	Gly	Ser	Lys	Gly	Val	Leu	Thr

Gly Ala Val <210> 4708 <211> 259 <212> PRT <213> Homo sapiens <400> 4708 Met Val Val Gln Gly Lys Arg Met Arg Lys Glu Thr Trp Gly Tyr Phe Cys Ser Lys Trp Asn Leu Leu Glu Leu Ala lle lle Leu Ala Ser Trp Ser Ala Leu Ala Val Phe Val Lys Arg Ala Val Leu Ala Glu Arg Asp Leu Gln Arg Cys Arg Asn His Arg Glu Glu Gly Ile Ser Phe Ser Glu Thr Ala Ala Ala Asp Ala Ala Leu Gly Tyr Ile Ile Val Phe Leu Val Leu Leu Ser Thr Val Lys Leu Trp His Leu Leu Arg Leu Asn Pro Lys Met Asn Met lle Thr Ala Ala Leu Arg Arg Ala Trp Gly Asp Ile Ser Gly Phe Ile Ile Val Ile Leu Thr Met Leu Leu Ala Tyr Ser Ile Ala Ser Asn Leu Ile Phe Gly Trp Lys Leu Arg Ser Tyr Lys Thr Leu Phe Asp Ala Ala Glu Thr Met Val Ser Leu Gln Leu Gly Ile Phe Asn Tyr

Glu Glu Val Leu Asp Tyr Ser Pro Val Leu Gly Ser Phe Leu Ile Gly

Ser Cys Ile Val Phe Met Thr Phe Val Val Leu Asn Leu Phe Ile Ser

 Val
 11e
 Leu
 Val
 Ala
 Phe
 Ser
 Glu
 Glu
 Glu
 Lys
 Tyr
 Glu
 Leu
 Ser

 Glu
 Glu
 Gly
 Glu
 Ile
 Val
 Asp
 Leu
 Leu
 Met
 Lys
 Ile
 Leu
 Ser
 Phe

 210
 21e
 21e
 21e
 Eu
 <td

<210> 4709

<211> 722

<212> PRT

<213> Homo sapiens

<400> 4709

 Met
 Lys
 Glu
 Val
 Leu
 Ser
 Thr
 Gly
 Gly
 Asn
 Thr
 Glu
 Val
 Ile

 1
 5
 10
 10
 15
 15

 His
 Thr
 Gly
 Thr
 Leu
 Gln
 Arg
 Tyr
 Gln
 Ser
 Tyr
 His
 Ile
 Gly
 Asp
 Phe

 Cys
 Phe
 Gln
 Glu
 Ile
 Glu
 Lys
 Glu
 Ile
 His
 Asp
 Ile
 Glu
 Phe
 Gln
 Cys

 Gln
 Glu
 Arg
 Asn
 Gly
 His
 Glu
 Ala
 Pro
 Met
 Thr
 Lys
 Ile
 Lys

50 55 60 Lys Leu Thr Gly Ser Thr Asp Gln His Asp His Arg His Ala Gly Asn 70 65 75 80 Lys Pro Ile Lys Asp Gln Leu Gly Ser Ser Phe Tyr Ser His Leu Pro 85 90 Glu Leu His Ile Ile Gln Ile Lys Gly Lys Ile Gly Asn Gln Phe Glu 105 Lys Ser Thr Ser Asp Ala Pro Ser Val Ser Thr Ser Gln Arg Ile Ser 115 120 Pro Arg Pro Gln Ile His Ile Ser Asn Asn Tyr Gly Asn Asn Ser Pro

	130					135					140				
Asn	Ser	Ser	Leu	Leu	Pro	Gln	Lys	Gln	Glu	Val	Tyr	Met	Arg	Glu	Lys
145					150					155					160
Ser	Phe	Gln	Cys	Asn	Glu	Ser	Gly	Lys	Ala	Phe	Asn	Cys	Ser	Ser	Leu
				165					170					175	
Leu	Arg	Lys	His	Gln	He	Pro	His	Leu	Gly	Asp	Lys	Gln	Tyr	Lys	Cys
			180					185					190		
Asp	Val	Cys	Gly	Lys	Leu	Phe	Asn	His	Lys	Gln	Tyr	Leu	Thr	Cys	His
		195					200					205			
Arg	Arg	Cys	His	Thr	Gly	Glu	Lys	Pro	Tyr	Lys	Cys	Asn	Glu	Cys	Gly
	210					215					220				
Lys	Ser	Phe	Ser	Gln	Val	Ser	Ser	Leu	Thr	Cys	His	Arg	Arg	Leu	His
225					230					235					240
Thr	Ala	Val	Lys	Ser	His	Lys	Cys	Asn	Glu	Cys	Gly	Lys	lle	Phe	Gly
		,		245					250					255	
Gln	Asn	Ser	Ala	Leu	Val	lle	His	Lys	Ala	lle	His	Thr	Gly	Glu	Lys
			260					265					270		
Pro	Tyr	Lys	Cys	Asn	Glu	Cys	Asp	Lys	Ala	Phe	Asn	Gln	Gln	Ser	Asn
		275					280					285			
Leu	Ala	Arg	His	Arg	Arg	Ile	His	Thr	Gly	Glu	Lys	Pro	Tyr	Lys	Cys
	290					295					300				
Glu	Glu	Cys	Asp	Lys	Val	Phe	Ser	Arg	Lys	Ser	Thr	Leu	Glu	Ser	His
305					310					315					320
Lys	Arg	lle	His	Thr	Gly	Glu	Lys	Pro	Tyr	Lys	Cys	Lys	Val	Cys	Asp
				325					330					335	
Thr	Ala	Phe	Thr	Trp	Asn	Ser	Gln	Leu	Ala	Arg	His	Lys	Arg	lle	His
			340					345					350		
Thr	Gly	Glu	Lys	Thr	Tyr	Lys	Cys	Asn	Glu	Cys	Gly	Lys	Thr	Phe	Ser
		355					360					365			
His	Lys	Ser	Ser	Leu	Val	Cys	His	His	Arg	Leu	His	Gly	Gly	Glu	Lys
	370					375					380				
Ser	Tyr	Lys	Cys	Lys		Cys	Asp	Lys	Ala		Ala	Trp	Asn	Ser	
385					390					395					400
Leu	Va]	Arg	His		Arg	He	His	Ser	Gly	Gly	Lys	Pro	Tyr	Lys	Cys
				405					410					415	
Asn	Glu	Cys	Gly	Lys	Thr	Phe	Gly	Gln	Asn	Ser	Asp	Leu	Leu	lle	llis

			420					425					430		
Lys	Ser	He	His	Thr	Gly	Glu	Gln	Pro	Tyr	Lys	Tyr	Glu	Glu	Cys	Glu
		435					440					445			
Lys	Val	Phe	Ser	Cys	Gly	Ser	Thr	Leu	Glu	Thr	His	Lys	He	11e	His
	450					455					460				
Thr	Gly	Glu	Lys	Pro	Tyr	Lys	Cys	Lys	Val	Cys	Asp	Lys	Ala	Phe	Ala
465					470					475					480
Cys	His	Ser	Tyr	Leu	Ala	Lys	His	Thr	Arg	lle	His	Ser	Gly	Glu	Lys
				485					490					495	
Pro	Tyr	Lys	Cys	Asn	Glu	Cys	Ser	Lys	Thr	Phe	His	Leu	Arg	Ser	Tyr
			500					505					510		
Leu	Ala	Ser	His	Arg	Arg	Val	His	Ser	Gly	Glu	Lys	Pro	Tyr	Lys	Cys
		515					520					525			
Asn	Glu	Cys	Ser	Lys	Thr	Phe	Ser	Gln	Arg	Ser	Tyr	Leu	His	Cys	His
	530					535					540				
Arg	Arg	Leu	His	Ser	Gly	Glu	Lys	Pro	Tyr	Lys	Cys	Asn	Glu	Cys	Gly
545					550					555					560
Lys	Thr	Phe	Ser	His	Lys	Pro	Ser	Leu	Val	His	His	Arg	Arg	Leu	His
				565					570					575	
Thr	Gly	Glu	Lys	Ser	Tyr	Lys	Cys	Thr	Val	Cys	Asp	Lys	Ala	Phe	Val
			580					585					590		
Arg	Asn	Ser	Tyr	Leu	Ala	Arg	His	Thr	Arg	He	His	Thr	Ala	Glu	Lys
		595					600					605			
Pro	Tyr	Lys	Cys	Asn	Glu	Cys	Gly	Lys	Ala	Phe	Asn	Gln	Gln	Ser	Gln
	610					615					620				
Leu	Ser	Leu	His	His	Arg	lle	His	Ala	Gly	Glu	Lys	Leu	Tyr	Lys	Cys
625					630					635					640
Glu	Thr	Cys	Asp		Val	Phe	Ser	Arg	Lys	Ser	His	Leu	Lys	Arg	His
				645					650					655	
Arg	Arg	He	His	Pro	Gly	Lys	Lys	Pro	Tyr	Lys	Cys	Lys		Cys	Asp
			660					665					670		
Lys	Thr		Gly	Ser	Asp	Ser		Leu	Lys	Gln	His		Gly	Leu	His
		675					680					685			
Thr		Glu	Lys	Pro	Tyr	Lys	Cys	Asn	Glu	Cys		Lys	Ala	Phe	Ser
	690	_				695					700				
Lvs	Gln	Ser	Thr	Leu	He	His	His	Gln	Ala	Val	His	Gly	Val	G] y	Lys

Leu Asp <210> 4710 <211> 422 <212> PRT <213> Homo sapiens <400> 4710 Met Phe Gln Thr Ala Trp Arg Gln Glu Pro Val Thr Phe Glu Asp Val Ala Val Tyr Phe Thr Gln Asn Glu Trp Ala Ser Leu Asp Ser Val Gln Arg Ala Leu Tyr Arg Glu Val Met Leu Glu Asn Tyr Ala Asn Val Ala Ser Leu Ala Phe Pro Phe Thr Thr Pro Val Leu Val Ser Gln Leu Glu Gln Gly Glu Leu Pro Trp Gly Leu Asp Pro Trp Glu Pro Met Gly Arg Glu Ala Leu Arg Gly Ile Cys Pro Gly Asp Glu Ala Arg Thr Glu Lys Glu Gly Leu Thr Pro Lys Asp His Val Ser Lys Glu Thr Glu Ser Phe Arg Leu Met Val Gly Gly Leu Pro Gly Asn Val Ser Gln His Leu Asp Phe Gly Ser Ser Leu Glu Gln Pro Gln Gly His Trp Ile Ile Lys Thr Lys Ser Lys Arg Arg His Phe Thr Asp Thr Ser Ala Arg His His Glu Ala Tyr Glu Val Lys Asn Gly Glu Lys Phe Glu Lys Leu Gly Lys Asn lle Ser Val Ser Thr Gln Leu Thr Thr Asn Gln Thr Asn Pro Ser Gly

Gln Ile Ser Tyr Glu Cys Gly Gln Cys Gly Arg Tyr Phe Ile Gln Met

		195					200					205			
Ala	Asp	Phe	His	Arg	His	Glu	Lys	Cys	His	Thr	Gly	Glu	Lys	Ser	Phe
	210					215					220				
Glu	Cys	Lys	Glu	Cys	Gly	Lys	Tyr	Phe	Arg	Tyr	Asn	Ser	Leu	Leu	lle
225					230					235					240
Arg	His	Gln	He	lle	His	Thr	Gly	Lys	Lys	Pro	Phe	Lys	Cys	Lys	Glu
				245					250					255	
Cys	Gly	Lys	Gly	Leu	Ser	Ser	Лѕр	Thr	Ala	Leu	lle	Gln	His	Gln	Arg
			260					265					270		
lle	His	Thr	Gly	Glu	Lys	Pro	Tyr	Glu	Cys	Lys	Glu	Cys	Gly	Lys	Ala
		275					280					285			
Phe	Ser	Ser	Ser	Ser	Val	Phe	Leu	Gln	His	Gln	Arg	Phe	His	Thr	Gly
	290					295					300				
Glu	Lys	Leu	Tyr	Glu	Cys	Asn	Glu	Cys	Trp	Lys	Thr	Phe	Ser	Cys	Ser
305					310					315					320
Ser	Ser	Phe	Thr	Val	His	Gln	Arg	Met	His	Thr	Gly	Glu	Lys	Pro	Tyr
				325					330					335	
Glu	Cys	Lys	Glu	Cys	Gly	Lys	Arg	Leu	Ser	Ser	Asn	Thr	Ala	Leu	Thr
			340					345					350		
Gln	His	Gln	Arg	He	His	Thr	Gly	Glu	Lys	Pro	Phe	Glu	Cys	Lys	Glu
		355					360					365			
Cys	Gly	Lys	Ala	Phe	Asn	Gln	Lys	He	Thr	Leu	lle	Gln	His	Gln	Arg
	370					375					380				
Val	His	Thr	Gly	Glu	Lys	Pro	Tyr	Glu	Cys	Lys	Val	Cys	G1 y	Lys	Thr
385					390					395					400
Phe	Ser	Trp	Cys	Gly	Arg	Phe	He	Leu	His	Gln	Lys	Leu	His	Thr	Gln
				405					410					415	
Lys	Thr	Pro	Val	Gln	Ala										
			420												

<210> 4711

<211> 446

<212> PRT

<213≻ Homo sapiens

<400	0> 47	711													
Met	Glu	Arg	lle	Pro	Val	Ser	Val	Asp	Phe	Trp	Val	Val	Cys	Cys	Ala
1				5					10					15	
Val	Leu	Lys	Cys	Asn	Pro	Gly	11e	Pro	Lys	Arg	Met	Ser	Thr	Leu	Cys
			20					25					30		
Phe	Gly	Phe	Ser	Asp	Glu	Phe	His	Pro	Phe	He	Glu	Ala	Leu	Leu	Pro
		35					40					45			
His	Val	Arg	Ala	He	Ala	Tyr	Thr	Trp	Phe	Asn	Leu	Gln	Ala	Gly	Lys
	50					55					60				
Arg	Lys	Tyr	Phe	Lys	Lys	His	Glu	Lys	Arg	Met	Ser	Lys	Asp	Glu	Glu
65					70					7.5					80
Arg	Ala	Val	Lys	Asp	Glu	Leu	Leu	Ser	Glu	Lys	Pro	Glu	lle	Lys	Gln
				85					90					95	
Lys	Trp	Ala	Ser	Arg	Leu	Leu	Ala	Lys	Leu	Arg	Lys	Asp	He	Arg	Gln
			100					105					110		
Glu	Tyr	Arg	Glu	Asp	Phe	Val	Leu	Thr	Val	Thr	Gly	Lys	Lys	His	Pro
		115					120					125			
Cys	Cys	Val	Leu	Ser	Asn	Pro	Asp	Gln	Lys	Gly	Lys	lle	Arg	Arg	Ile
	130					135					140				
Asp	Cys	Leu	Arg	Gln	Ala	Asp	Lys	Va]	Trp	Arg	Leu	Asp	Leu	Val	Met
145					150					155					160
Val	lle	Leu	Phe	Lys	Gly	11e	Pro	Leu	Glu	Ser	Thr	Asp	Gly	Glu	Arg
				165					170					175	
Leu	Met	Lys	Ser	Pro	His	Cys	Thr	Asn	Pro	Ala	Leu	Cys	Val	Gln	Pro
			180					185					190		
His	His	Ile	Thr	Va]	Ser	Val	Lys	Glu	Leu	Asp	Leu	Phe	Leu	Ala	Tyr
		195					200					205			
Tyr		Gln	Glu	Gln	Asp	Ser	G1 y	Gln	Ser	G1 y	Ser	Pro	Ser	His	Ser
	210					215					220				
Asp	Pro	Ala	Lys	Asn	Pro	Pro	Gly	Tyr	Leu	Glu	Asp	Ser	Phe	Val	Lys
225					230					235					240
Ser	Gly	Val	Phe		Val	Ser	G] u	Leu		Arg	Val	Ser	Arg	Thr	Pro
				245					250					255	
Ile	Thr	G1n		Thr	G1 y	Val	Asn		Pro	He	Gly	Glu	He	Pro	Ser
			260					265					270		
Gln	Pro	Tyr	Tyr	His	Asp	Met	Asn	Ser	Gly	Val	Asn	Leu	Gln	Arg	Ser

Leu Ser Ser Pro Pro Ser Ser Lys Arg Pro Lys Thr lle Ser lle Asp Glu Asn Met Glu Pro Ser Pro Thr Gly Asp Phe Tyr Pro Ser Pro Ser Ser Pro Ala Ala Gly Ser Arg Thr Trp His Glu Arg Asp Gln Asp Met Ser Ser Pro Thr Thr Met Lys Lys Pro Glu Lys Pro Leu Phe Ser Ser Ala Ser Pro Gln Asp Ser Ser Pro Arg Leu Ser Thr Phe Pro Gln His His His Pro Gly 11e Pro Gly Val Ala His Ser Val 11e Ser Thr Arg Thr Pro Pro Pro Pro Ser Pro Leu Pro Phe Pro Thr Gln Ala 11e Leu Pro Pro Ala Pro Ser Ser Tyr Phe Ser His Pro Thr lle Arg Tyr Pro Pro His Leu Asn Pro Gln Asp Thr Leu Lys Asn Tyr Val Pro Ser Tyr Asp Pro Ser Ser Pro Gln Thr Ser Gln Ser Trp Tyr Leu Gly

<210> 4712

<211> 488

<212> PRT

<213> Homo sapiens

<400> 4712

	50					55					60				
Val	Asn	Gly	Leu	Tyr	Ser	Ser	Ser	Asp	Asp	Val	He	Glu	Leu	Thr	Pro
65					70					75					80
Ser	Asn	Phe	Asn	Arg	Glu	Val	He	Gln	Ser	Лѕр	Ser	Leu	Trp	Leu	Val
				85					90					95	
Glu	Phe	Tyr	Ala	Pro	Trp	Cys	Gly	His	Cys	Gln	Arg	Leu	Thr	Pro	Glu
			100					105					110		
Trp	Lys	Lys	Ala	Ala	Thr	Ala	Leu	Lys	Asp	Val	Val	Lys	Val	Gly	Ala
		115					120					125			
Val	Asp	Ala	Asp	Lys	His	His	Ser	Leu	G1 y	Gly	Gln	Tyr	Gly	Val	Gln
	130					135					140				
Gly	Phe	Pro	Thr	lle	Lys	He	Phe	Gly	Ser	Asn	Lys	Asn	Arg	Pro	Glu
145					150					155					160
Asp	Tyr	Gln	Gly	Gly	Arg	Thr	Gly	Glu	Ala	He	Val	Asp	Ala	Ala	Leu
				165					170					175	
Ser	Ala	Leu	Arg	Gln	Leu	Val	Lys	Asp	Arg	Leu	Gly	Gly	Arg	Ser	G1 y
			180					185					190		
Gly	Tyr	Ser	Ser	Gly	Lys	Gln	Gly	Arg	Ser	Asp	Ser	Ser	Ser	Lys	Lys
		195					200					205			
Asp	Val	lle	Glu	Leu	Thr	Asp	Asp	Ser	Phe	Asp	Lys	Asn	Val	Leu	Asp
	210					215					220				
Ser	Glu	Asp	Val	Trp	Met	Val	Glu	Phe	Tyr	Ala	Pro	Trp	Cys	Gly	His
225					230					235			-		240
Cys	Lys	Asn	Leu	Glu	Pro	Glu	Trp	Ala	Ala	Ala	Ala	Ser	Glu	Val	Lys
				245					250					255	
Glu	Gln	Thr	Lys	Gly	Arg	Val	Lys	Leu	Ala	Ala	Val	Asp	Ala	Thr	Va]
			260					265					270		
Asn	Gln		Leu	Ala	Ser	Arg		G1 y	lle	Arg	Gly		Pro	Thr	He
		275					280					285			
Lys		Phe	Gln	Lys	Gly		Ser	Pro	Val	Asp		Asp	Gly	Gly	Arg
	290					295					300				
	Arg	Ser	Asp	He		Ser	Arg	Ala	Leu		Leu	Phe	Ser	Asp	
305					310					315					320
Ala	Pro	Pro	Pro		Leu	Leu	Glu	He		Asn	Glu	Asp	He		Lys
				325				_	330					335	
Arg	Thr	Cys	Glu	Glu	His	GIn	Leu	Cys	Val	Val	Ala	Val	Leu	Pro	His

			340					345					350		
He	Leu	Asp	Thr	Gly	Ala	Ala	Gly	Arg	Asn	Ser	Tyr	Leu	Glu	Val	Leu
		355					360					365			
Leu	Lys	Leu	Ala	Asp	Lys	Tyr	Lys	Lys	Lys	Met	Trp	Gly	Trp	Leu	Trp
	370					375					380				
Thr	Glu	Ala	G1 y	Ala	Gln	Ser	Glu	Leu	Glu	Thr	Ala	Leu	Gly	lle	Gly
385					390					395					400
Gly	Phe	Gly	Tyr	Pro	Λla	Met	Ala	Ala	lle	Asn	Ala	Arg	Lys	Met	Lys
				405					410					415	
Phe	Ala	Leu	Leu	Lys	Gly	Ser	Phe	Ser	Glu	Gln	Gly	He	Asn	Glu	Phe
			420					425					430		
Leu	Arg	Glu	Leu	Ser	Phe	Gly	Arg	Gly	Ser	Thr	Ala	Pro	Val	Gly	Gly
		435					440					445			
Gly	Ala	Phe	Pro	Thr	He	Val	Glu	Arg	Glu	Pro	Trp	Asp	Gly	Arg	Asp
	450					455					460				
Gly	Glu	Leu	Pro	Val	Glu	Asp	Asp	lle	Asp	Leu	Ser	Asp	Val	Glu	Leu
465					470					475					480
Asp	Asp	Leu	Gly	Lys	Asp	Glu	Leu								
				485											
<210)> 4°	713													
<21	1> 3	18													
<212	2> PI	RT													
<213	3> H	omo :	sapie	ens											
<400)> 47	713													
Met	Val	Leu	Ser	Ser	Gly	Pro	Gln	Trp	Cys	G1 y	Ser	Gln	Glu	Leu	Trp
1				5					10					15	
Phe	Gly	Lys	Thr	Cys	Glu	Glu	Lys	Ser	Arg	Leu	Gly	Arg	Trp	Pro	Gly
			20					25					30		
Tyr	Leu	Asn	Gly	G1 y	Arg	Met	G1u	Ser	Ser	Thr	Asn	Asp	He	Пе	Glu
		35					40					45			
Val	lle	Val	Lys	Asp	Glu	Met	He	Ser	Val	Glu	Glu	Ser	Ser	Gly	Asn

Thr	Asp	Val	Asn	Asn	Leu	Leu	Gly	lle	His	His	Lys	Ile	Leu	Asn	Glu
65					70					75					80
Gln	Ile	Phe	Tyr	He	Cys	Glu	Glu	Cys	Gly	Lys	Cys	Phe	Asp	Gln	Asn
				85					90					95	
Glu	Asp	Phe	Asp	Gln	His	Gln	Lys	Thr	His	Asn	Gly	Glu	Lys	Val	Tyr
			100					105					110		
Gly	Cys	Lys	Glu	Cys	Gly	Lys	Ala	Phe	Ser	Phe	Arg	Ser	His	Cys	He
		115					120					125			
Ala	His	Gln	Arg	Ile	His	Ser	Gly	Val	Lys	Pro	Tyr	Glu	Cys	Gln	Glu
	130					135					140				
Cys	Ala	Lys	Ala	Phe	Val	Trp	Lys	Ser	Asn	Leu	Ile	Arg	His	Gln	Arg
145					150					155					160
He	His	Thr	Gly	Glu	Lys	Pro	Phe	Glu	Cys	Lys	Glu	Cys	Gly	Lys	Gly
				165					170					175	
Phe	Ser	Gln	Asn	Thr	Ser	Leu	Thr	Gln	His	Gln	Arg	lle	His	Thr	Gly
			180					185					190		
Glu	Lys	Pro	Tyr	Thr	Cys	Lys	Glu	Cys	Gly	Lys	Ser	Phe	Thr	Arg	Asn
		195					200					205			
Pro	Ala	Leu	Leu	Arg	His	Gln	Arg	Met	His	Thr	Gly	Glu	Lys	Pro	Tyr
	210					215					220				
Glu	Cys	Lys	Asp	Cys	G1y	Lys	Gly	Phe	Met	Trp	Asn	Ser	Asp	Leu	Ser
225					230					235					240
Gln	His	Gln	Arg	Val	His	Thr _.	Gly	Asp	Lys	Pro	His	Glu	Cys	Thr	Asp
				245					250					255	
Cys	Gly	Lys	Ser	Phe	Phe	Cys	Lys	Ala	His	Leu	He	Arg	His	Gln	Arg
			260					265					270		
lle	His	Thr	Gly	Glu	Arg	Pro	Tyr	Lys	Cys	Asn	Asp	Cys	Gly	Lys	Ala
		275					280					285			
Phe	Ser	Gln	Asn	Ser	Val	Leu	He	Lys	His	Gln	Arg	Arg	His	Ala	Arg
	290					295					300				
Лsp	Lys	Pro	Tyr	Asn	Cys	Gln	lle	Ser	llis	Leu	Leu	Glu	His		
305					310					315					

<210> 4714 <211> 743

<212	2> PF	RT.													
<213	3> Ho	omo s	sapie	ens											
<400)> 47	714													
Met	He	Ser	Gln	Phe	Phe	He	Leu	Ser	Ser	Lys	Gly	Asp	Pro	Leu	He
1				5					10					15	
Tyr	Lys	Asp	Phe	Arg	Gly	Asp	Ser	Gly	Gly	Arg	Asp	Val	Ala	Glu	Leu
			20					25					30		
Phe	Tyr	Arg	Lys	Leu	Thr	Gly	Leu	Pro	Gly	Asp	Glu	Ser	Pro	Val	Val
		35					40					45			
Met	Asp	Tyr	Gly	Tyr	Val	Gln	Thr	Thr	Ser	Thr	Glu	Met	Leu	Arg	Asn
	50					55					60				
Phe	11e	Gln	Thr	Glu	Ala	Va]	Val	Ser	Lys	Pro	Phe	Ser	Leu	Phe	Asp
65					70					75					80
Leu	Ser	Ser	Val	Gly	Leu	Phe	Gly	Ala	Glu	Thr	Gln	Gln	Ser	Lys	Val
				85					90					95	
Ala	Pro	Ser	Ser	Ala	Ala	Ser	Arg	Pro	Val	Leu	Ser	Ser	Arg	Ser	Asp
			100					105					110		
Gln	Ser	Gln	Lys	Asn	Glu	Val	Phe	Leu	Asp	Val	Val	Glu	Arg	Leu	Ser
		115					120					125			
Val	Leu	Ile	Ala	Ser	Asn	Gly	Ser	Leu	Leu	Lys	Val	Asp	Val	Gln	Gly
	130					135					140				
Glu	lle	Arg	Leu	Lys	Ser	Phe	Leu	Pro	Ser	Gly	Ser	Glu	Met	Arg	11e
145					150					155					160
Gly	Leu	Thr	Glu	Glu	Phe	Cys	Val	Gly	Lys	Ser	Glu	Leu	Arg	Gly	Tyr
				165					170					175	
Gly	Pro	Gly	lle	Arg	Val	Asp	Glu	Val	Ser	Phe	His	Ser	Ser	Val	Asn
			180					185					190		
Leu	Asp	Glu	Phe	Glu	Ser	His	Arg	lle	Leu	Arg	Leu	Gln	Pro	Pro	Gln
		195					200					205			
Gly	Glu	Leu	Thr	Val	Met	Arg	Tyr	Gln	Leu	Ser	Asp	Asp	Leu	Pro	Ser
	210					215					220				
Pro	Leu	Pro	Phe	Arg	Leu	Phe	Pro	Ser	Val	Gln	Trp	Asp	Arg	Gly	Ser

Gly Arg Leu Gln Val Tyr Leu Lys Leu Arg Cys Asp Leu Leu Ser Lys

Ser	Gln	Ala	Leu	Asn	Val	Arg	Leu	His	Leu	Pro	Leu	Pro	Arg	Gly	Val
			260					265					270		
Val	Ser	Leu	Ser	Gln	Glu	Leu	Ser	Ser	Pro	Glu	Gln	Lys	Ala	Glu	Leu
		275					280					285			
Λla	Glu	Gly	Ala	Leu	Arg	Trp	Asp	Leu	Pro	Arg	Val	Gln	Gly	Gly	Ser
	290					295					300				
Gln	Leu	Ser	Gly	Leu	Phe	Gln	Ser	Arg	Lys	Gly	Ala	Asp	Leu	Asp	Arg
305					310					315					320
Glu	Lys	Lys	Ala	Ala	Glu	Cys	Lys	Val	Asp	Ser	Ile	Gly	Ser	G1y	Arg
				325					330					335	
Ala	He	Pro	lle	Lys	Gln	Gly	Ile	Leu	Leu	Lys	Arg	Ser	Gly	Lys	Ser
			340					345					350		
Leu	Asn	Lys	Glu	Trp	Lys	Lys	Lys	Tyr	Val	Thr	Leu	Cys	Лsp	Asn	Gly
		355					360					365			
Leu	Leu	Thr	Tyr	His	Pro	Ser	Leu	His	Asp	Tyr	Met	Gln	Asn	Пе	His
	370					375					380				
Gly	Lys	Glu	lle	Asp	Leu	Leu	Arg	Thr	Thr	Val	Lys	Val	Pro	Gly	Lys
385					390					395					400
Arg	Leu	Pro	Arg	Ala	Thr	Pro	Ala	Thr	Ala	Pro	Gly	Thr	Ser	Pro	Arg
				405					410					415	
Ala	Asn	Gly	Leu	Ser	Val	Glu	Arg	Ser	Asn	Thr	Gln	Leu	Gly	Gly	Gly
			420					425					430		
Thr	Glu	Ala	Glu	Glu	Ser	Phe	Glu	Phe	Val	Val	Val	Ser	Leu	Thr	Gly
		435					440					445			
Gln	Thr	Trp	His	Phe	Glu	Ala	Ser	Thr	Ala	Glu	Glu	Arg	Glu	Leu	Trp
	450					455					460				
Val	Gln	Ser	Val	Gln	Ala	Gln	He	Leu	Ala	Ser	Leu	Gln	Gly	Cys	Arg
465					470					475					480
Ser	Ala	Lys	Asp	Lys	Thr	Arg	Leu	Gly	Asn	Gln	Asn	Ala	Ala	Leu	Ala
				485					490					495	
Val	Gln	Ala	Val	Arg	Thr	Val	Arg	Gly	Asn	Ser	Phe	Cys	lle	Λsp	Cys
			500					505					510		
Asp	Ala	Pro	Asn	Pro	Asp	Trp	Ala	Ser	Leu	Asn	Leu	Gly	Ala	Leu	Met
		515					520					525			
Cys	lle	Glu	Cys	Ser	Gly	lle	His	Arg	His	Leu	Gly	Ala	His	Leu	Ser
	530					535					540				

A	$V_{\rm C}$ 1	A 200	S 0 22	1	A 615	Lav	Acr	Acr	Tre	Dro	Dra	C1	1 600	1	۸1.
	vai	Arg	ser	Leu		Leu	Asp	ASP	ırp		rro	oru	Leu	Leu	
545					550					555					560
Val	Met	Thr	Ala	Met	Gly	Asn	Ala	Leu	Ala	Asn	Ser	Val	Trp	Glu	Gly
				565					570					575	
Ala	Leu	Gly	Gly	Tyr	Ser	Lys	Pro	G1 y	Pro	Asp	Ala	Cys	Arg	Glu	Glu
			580					585					590		
Lys	Glu	Arg	Trp	lle	Arg	Ala	Lys	Tyr	Glu	Gln	Lys	Leu	Phe	Leu	Ala
		595					600					605			•
Pro	Leu	Pro	Ser	Ser	Asp	Val	Pro	Leu	Gly	Gln	Gln	Leu	Leu	Arg	Ala
	610					615					620				
Val	Va]	Glu	Asp	Asp	Leu	Arg	Leu	Leu	Val	Met	Leu	Leu	Ala	His	Gly
625					630					635					640
	Lys	Glu	Glu	Va1		Glu	Thr	Tyr	Gly	Asp	Gly	Asp	Gly	Arg	Thr
				645					650					655	
Ala	Leu	His	Leu	Ser	Ser	Ala	Met	Ála		Val	Val	Phe	Thr		Leu
			660					665					670		
Lau	Ha	Trn		Glv	Val	Asp	Val		Sar	Ara	Asn	Ala		Glv	ادم ا
Leu	116		1 9 1	Gry	vaı	nsp		AI g	361	ni g	nsp		nı g	Oly	Leu
	_	675		_			680			_		685	_		
Thr		Leu	Ala	Tyr	Ala	Arg	Arg	Ala	Gly	Ser	Gln	Glu	Cys	Ala	Asp
	690					695				•	700				
He	Leu	lle	Gln	His	Gly	Cys	Pro	Gly	Glu	Gly	Cys	Gly	Leu	Ala	Pro
705					710					715					720
Thr	Pro	Asn	Arg	Glu	Pro	Ala	Asn	Gly	Thr	Asn	Pro	Ser	Ala	Glu	Leu
				725					730					735	
His	Arg	Ser	Pro	Ser	Leu	Leu									
			740												

<210> 4715

<211> 141

<212> PRT

<213> Homo sapiens

<400> 4715

Met Gln Ile Gln Glu Ile Gln Arg Thr Pro Leu Arg Tyr Ser Lys Arg 1 5 10 15 Arg Ser Thr Pro Arg Pro Leu Ile Ile Gly Phe Ser Lys Val Glu Leu 25 Leu Arg Ala Ala Arg Glu Lys Gly Gln Val Thr Tyr Lys Gly Lys Pro 40 45 lle Arg Pro Thr Ala Asp Phe Ser Ala Glu Pro Leu Gln Ala Arg Arg 55 60 Asp Cys Gly Pro Ile Ser Asn Ile Leu Lys Gly Lys Asn Phe Gln Pro 70 75 Arg Ile Ser Tyr Pro Ala Lys Leu Ile Ser Phe Ile Ser Glu Gly Glu 85 90 Ile Lys Ser Phe Pro Asp Lys Gln Met Leu Arg Asp Phe Val Thr Thr 105 Arg Pro Ala Leu Gln Glu Leu Leu Lys Glu Ala Leu Asn Met Glu Arg 115 120 Lys Asn Gln Tyr Gln Pro Leu Gln Lys His Thr Lys 11e 130 135 140

<210> 4716

<211> 715

<212> PRT

<213> Homo sapiens

<400> 4716

Met Ala Thr Phe Met Asp Pro Gly Val Phe Pro Arg Ala Asp Glu Asp 1 5 10 15

Glu Asp Lys Glu Asp Asp Phe Arg Ala Pro Leu Tyr Lys Asn Val Asp
20 25 30

Val Arg Gly 11e Gln Val Arg Met Lys Trp Cys Ala Thr Cys His Phe 35 40 45

Tyr Arg Pro Pro Arg Cys Ser His Cys Ser Val Cys Asp Asn Cys Val
50 55 60

Glu Asp Phe Asp His His Cys Pro Trp Val Asn Asn Cys lle Gly Arg
65 70 75 80

Arg Asn Tyr Arg Tyr Phe Phe Leu Phe Leu Leu Ser Leu Ser Ala His

85

Met	Val	Gly	Val	Val	Ala	Phe	Gly	Leu	Val	Tyr	Val	Leu	Asn	His	Ala
		_	100					105					110		
Glu	Gly	Leu	Gly	Ala	Ala	His	Thr	Thr	Пe	Thr	Met	Ala	Val	Met	Cys
		115					120					125			
Val	Ala	Gly	Leu	Phe	Phe	lle	Pro	Val	11e	Gly	Leu	Thr	Gly	Phe	His
	130					135					140				
Val	Val	Leu	Val	Thr	Arg	Gly	Arg	Thr	Thr	Asn	Glu	Gln	Val	Thr	Gly
145					150					155					160
Lys	Phe	Arg	Gly	Gly	Val	Asn	Pro	Phe	Thr	Arg	Gly	Cys	Cys	Gly	Asn
				165					170					175	
Val	Glu	His	Val	Leu	Cys	Ser	Pro	Leu	Ala	Pro	Arg	Tyr	Val	Val	Glu
			180					185					190		
Pro	Pro	Arg	Leu	Pro	Leu	Ala	Val	Ser	Leu	Lys	Pro	Pro	Phe	Leu	Arg
		195					200					205			
Pro	Glu	Leu	Leu	Asp	Arg	Ala	Ala	Pro	Leu	Lys	Val	Lys	Leu	Ser	Asp
	210					215					220				
Asn	Gly	Leu	Lys	Ala	Gly	Leu	Gly	Arg	Ser	Lys	Ser	Lys	Gly	Ser	
225					230					235					240
Asp	Arg	Leu	Asp		Lys	Pro	Leu	Asp		Gly	Pro	Pro	Leu	Pro	Pro
				245					250					255	
Lys	He	Glu		Gly	Thr	Phe	Ser		Asp	Leu	Gln	Thr		Arg	Pro
			260	_				265					270		
Gly	Ser		Glu	Ser	Ala	Leu		Val	Gln	Arg	Thr		Pro	Pro	Thr
		275			121		280		73.1		6 01	285	15		
Pro		Met	Tyr	Lys	Phe		Pro	Ala	Phe	Pro		Gly	Pro	Lys	Val
10	290 Di	0	61	D	C1	295	C1	V 1	n	C1	300		C	,	T.I
	Phe	Cys	Gly	Pro		Glu	GIn	vai	Pro		Pro	Asp	Ser	Leu	
305	C1 .	Δ.	Α	C	310	A	C	1	A	315	17 - 1	C	C1	D	320
Leu	61 y	Asp	Asp		116	Arg	ser	Leu		rne	vai	Ser	GIU	Pro	Ser
1	۸	1	D	325	Т	C1	Dana	C1	330	Lan	114	110	110	335	Dana
Leu	ASP	Leu		ASP	Tyr	GIY	110		GIY	Leu	nis	V19	350	Tyr	110
Dava	C	Dua	340 Bras	l a	Com	110	Can	345	۸1.	Dha	C	C1		Lau	Λ
r 10	ser	355	110	Leu	Set.	nıa	360	nsp	MIA	гие	per.	365	nia	Leu	wi.8
Sor	Lou		lou	lve	Δla	Sor		Arg	Ara	Glv	610		Hic	Val	Ala
Sel.	370	261	Leu	LYS	ліа	375	261	мв	лл	оту	380	nsp	1112	101	nia

Leu	GIn	Pro	Leu	Arg	Ser	Glu	Gly	61 y	Pro	Pro	Ihr	Pro	HIS	Arg	5er
385					390					395					400
He	Phe	Ala	Pro	His	Ala	Leu	Pro	Asn	Arg	Asn	Gly	Ser	Leu	Ser	Tyr
				405					410					415	
Asp	Ser	Leu	Leu	Asn	Pro	Gly	Ser	Pro	Gly	Gly	His	Ala	Cys	Pro	Ala
			420					425					430		
llis	Pro	Ala	Val	G]y	Val	Ala	Gly	Tyr	His	Ser	Pro	Tyr	Leu	llis	Pro
		435					440					445			
Gly	Ala	Thr	Gly	Asp	Pro	Pro	Arg	Pro	Leu	Pro	Arg	Ser	Phe	Ser	Pro
	450					455					460				
Val	Leu	Gly	Pro	Arg	Pro	Arg	Glu	Pro	Ser	Pro	Val	Arg	Tyr	Asp	Asn
465					470					475					480
Leu	Ser	Arg	Thr	lle	Met	Ala	Ser	lle	Gln	Glu	Arg	Lys	Аsp	Arg	Glu
				485					490					495	
Glu	Arg	Glu	Arg	Leu	Leu	Arg	Ser	Gln	Ala	Asp	Ser	Leu	Phe	Gly	Asp
			500					505					510		
Ser	Gly	Val	Tyr	Asp	Ala	Pro	Ser	Ser	Tyr	Ser	Leu	Gln	Gln	Ala	Ser
		515					520					525			
Val	Leu	Ser	Glu	Gly	Pro	Arg	Gly	Pro	Ala	Leu	Arg	Tyr	Gly	Ser	Arg
	530					535					540				
Asp	Asp	Leu	Val	Ala	Gly	Pro	Gly	Phe	Gly	G1 y	Ala	Arg	Asn	Pro	Ala
545					550					555					560
Leu	Gln	Thr	Ser	Leu	Ser	Ser	Leu	Ser	Ser	Ser	Val	Ser	Arg	Ala	Pro
				565					570					575	
Arg	Thr	Ser	Ser	Ser	Ser	Leu	Gln		Asp	Gln	Ala	Ser		Asn	Ala
			580					585					590		
Pro	Gly		Arg	Pro	Ser	Ser		Ser	His	Arg	Ser		Ala	Arg	G1n
		595					600					605	_		
Gly		Pro	Ser	Pro	Pro		Thr	Pro	His	Ser		Ser	Tyr	Ala	G1 y
	610					615				_	620		***		
	Lys	Ala	Val	Ala		lle	His	Thr	Asp		Pro	Glu	Pro	Pro	
625	_				630					635	_			0.7	640
Ser	Leu	Thr	Val		Arg	Gly	Arg	He		Thr	Cys	Thir	Arg	Gly	Trp
0.7			63	645	ъ	m.	., .		650	0.				655	
Gly	Arg	Arg	-	GIn	Pro	Trp	Val		Pro	Gly	Leu	His		Cys	His
			660					665					670		

Leu Gly Arg Pro Glu Asp Arg Pro Pro Leu Arg Ala Pro Trp Ser Gln Ala Ala Gly Ala Pro Pro Arg Gly Ala Met Cys Arg Leu His Leu Ala Ala Ser Ser Leu Phe Pro Ser Leu Ser Gly Pro

<210> 4717

<211> 851

<212> PRT

<213> Homo sapiens

<400> 4717 Met Ser Val Ser Phe His Thr His Thr Lys Glu Leu Trp Thr Trp Met Glu Asp Leu Gln Lys Glu Met Leu Glu Asp Val Cys Ala Asp Ser Val Asp Ala Val Gln Glu Leu Ile Lys Gln Phe Gln Gln Gln Gln Thr Ala Thr Leu Asp Ala Thr Leu Asn Val Ile Lys Glu Gly Glu Asp Leu Ile Gln Gln Leu Arg Asp Ser Ala Val Ser Asn Asn Lys Thr Pro His Ser Ser Ser lle Ser His Ile Glu Ser Val Leu Gln Gln Leu Asp Asp Ala Gln Val Gln Met Glu Glu Leu Phe His Glu Arg Lys Ile Lys Leu Asp lle Phe Leu Gln Leu Arg lle Phe Glu Gln Tyr Thr lle Glu Val Thr Ala Glu Leu Asp Ala Trp Asn Glu Asp Leu Leu Arg Gln Met Asn Asp

Phe Asn Thr Glu Asp Leu Thr Leu Ala Glu Gln Arg Leu Gln Arg His

Thr Glu Arg Lys Leu Ala Met Asn Asn Met Thr Phe Glu Val Ile Gln

Gln	Gly	Gln	Asp	Leu	His	Gln	Tyr	Ile	Thr	Glu	Val	Gln	Ala	Ser	Gly
			180					185					190		
Пе	Glu	Leu	He	Cys	Glu	Lys	Asp	Ile	Asp	Leu	Ala	Ala	Gln	Val	Gln
		195					200					205			
Glu	Leu	Leu	Glu	Phe	Leu	His	Glu	Lys	Gln	His	Glu	Leu	Glu	Leu	Asn
	210					215					220				
Ala	Glu	Gln	Thr	His	Lys	Arg	Leu	Glu	Gln	Cys	Leu	Gln	Leu	Arg	His
225					230					235					240
Leu	Gln	Ala	Glu	Val	Lys	Gln	Val	Leu	Gly	Trp	Ile	Arg	Asn	Gly	Glu
				245					250					255	
Ser	Met	Leu	Asn	Ala	Ser	Leu	Val	Asn	Ala	Ser	Ser	Leu	Ser	Glu	Ala
			260					265					270		
Glu	Gln	Leu	Gln	Arg	Glu	His	Glu	Gln	Phe	Gln	Leu	Ala	He	Glu	Ser
		275					280					285			
Leu	Phe	llis	Ala	Thr	Ser	Leu	Gln	Lys	Thr	His	Gln	Ser	Ala	Leu	Gln
	290					295					300				
Val	Gln	Gln	Lys	Ala	Glu	Val	Leu	Leu	Gln	Ala	Gly	His	Tyr	Asp	Ala
305					310					315					320
Asp	Ala	Ile	Arg	Glu	Cys	Ala	Glu	Lys	Val	Ala	Leu	His	Trp	Gln	Gln
				325					330					335	
Leu	Met	Leu	Lys	Met	Glu	Asp	Arg	Leu	Lys	Leu	Val	Asn	Ala	Ser	Val
			340					345					350		
Ala	Phe	Tyr	Lys	Thr	Ser	Glu	Gln	Val	Cys	Ser	Val	Leu	Glu	Ser	Leu
		355					360					365			
Glu	Gln	Glu	Tyr	Arg	Arg	Asp	Glu	Asp	Trp	Cys	Gly	Gly	Arg	Asp	Lys
	370					375					380				
Leu	Gly	Pro	Ala	Ala	Glu	lle	Asp	His	Val	lle	Pro	Leu	lle	Ser	Lys
385					390					395					400
llis	Leu	Glu	Gln	Lys	Glu	Ala	Phe	Leu	Lys	Ala	Cys	Thr	Leu	Ala	Arg
				405					410					415	
Arg	Asn	Ala	Glu	Va]	Phe	Leu	Lys	Tyr	He	His	Arg	Asn	Asn	Val	Ser
			420					425					430		
Met	Pro	Ser	Val	Ala	Ser	His	Thr	Arg	Gly	Pro	Glu	Gln	Gln	Val	Lys
		435					440					445			
Ala	He	Leu	Ser	Glu	Leu	Leu	Gln	Arg	Glu	Asn	Arg	Val	Leu	His	Phe
	450					455					460				

Trp	Thr	Leu	Lys	Lys	Arg	Arg	Leu	Asp	Gln	Cys	Gln	Gln	Tyr	Val	Val
465					470					475					480
Phe	Glu	Arg	Ser	Ala	Lys	Gln	Ala	Leu	Asp	Trp	He	Gln	Glu	Thr	Gly
				485					490					495	
Glu	Phe	Tyr	Leu	Ser	Thr	His	Thr	Ser	Thr	Gly	Glu	Thr	Thr	Glu	Glu
			500					505					510		
Thr	Gln	Glu	Leu	Leu	Lys	Glu	Tyr	Gly	Glu	Phe	Arg	Val	Pro	Ala	Lys
		515					520					525			
Gln	Thr	Lys	Glu	Lys	Val	Lys	Leu	Leu	He	Gln	Leu	Ala	Asp	Ser	Phe
	530					535					540				
Val	Glu	Lys	Gly	His	lle	His	Ala	Thr	Glu	Ile	Arg	Lys	Trp	Val	Thr
545					550					555					560
Thr	Va]	Asp	Lys	His	Tyr	Arg	Asp	Phe	Ser	Leu	Arg	Met	Gly	Lys	Tyr
				565					570					575	
Arg	Tyr	Ser	Leu	Glu	Lys	Ala	Leu	Gly	Val	Asn	Thr	Glu	Asp	Asn	Lys
			580					585					590		
Asp	Leu		Leu	Asp	lle	lle		Ala	Ser	Leu	Ser	Asp	Arg	Glu	Val
		595					600					605			
Lys		Arg	Asp	Ala	Asn		Glu	Val	Asn	Glu		Lys	Arg	Lys	Ser
	610					615					620				
	Arg	Lys	Lys	Glu		lle	Met	Ala	Glu		Leu	Gln	Thr	G] u	
625	_				630			_		635					640
Ala	Tyr	Val	Arg		Leu	His	Glu	Cys		Glu	Thr	Tyr	Leu		Glu
	m)	0	0.1	645	0.1	0.1		D	650	0.1		,		655	6.1
Met	Thr	Ser	Gly	Val	Glu	Glu	He		Pro	Gly	11e	Leu		Lys	Glu
11.7	т1.	71.	660	C1		11.	C1	665	11.	т	۸	Dl	670	۸	Λ
HIS	11e		Phe	Gly	Asn	116		GIU	116	lyr	Asp		HIS	ASN	Asn
11.	Dl. a	675	1	C1	1	C1	680	т	C1	C1	Lau	685	C1	Aon	V o 1
116	690	Leu	Lys	Gju	Leu	695	Lys	Tyr	GIU	OIII	700	110	Gju	кѕр	vai
Gl v		Cvc	Phe	Vol	The		Ala	Acn	Lve	Pho		Mot	Tyr	Val	Thr
705	1115	Cys	THE	vai	710	11 b	MIA	nsp	Lys	715	nto	sie t	1 y 1	101	720
	Cve	lve	Asn	Lve		Asn	Ser	Aen	Gln		He	Len	Glu	His	
۱. و ۱	0,3	r. 3 13	11311	725	110	чыр	561	41.211	730	150 U	.1 .1 .	,,,u	Ç, i u	735	
Glv	Thr	Phe	Phe		Glu	He	Gln	Gln		His	Glv	Leu	Ala		Ser
013		1 ,10	740	op	Olu		OTH	745	8	1,13	01	.,,,,,	750		~ ~ 1

Ile Ser Ser Tyr Leu Ile Lys Pro Val Gln Arg Ile Thr Lys Tyr Gln Leu Leu Leu Lys Glu Leu Leu Thr Cys Cys Glu Glu Gly Lys Gly Glu Leu Lys Asp Gly Leu Glu Val Met Leu Ser Val Pro Lys Lys Ala Asn Asp Ala Met His Val Ser Met Leu Glu Gly Ser Cys Pro Pro Ser Thr Gly Glu Ala Ser Ser Leu Pro Arg His Gly Gly Ala Cys Ile Met Gly Gly Lys Trp His Glu Val Arg Gln Gly Ala Arg Leu Glu Glu Arg Arg Asn Asp Lys

<210> 4718

<211> 312

<212> PRT

<213> Homo sapiens

<400> 4718

Met lle Ala Val Gly Ser Met Asp Tyr Gly Leu Trp Gln Leu Phe Cys Thr Leu Glu Leu Pro Leu 11e Pro 11e Leu Ala Val Met Glu Ser His Ala Ile Gln Val Asn Lys Glu Glu Met Glu Lys Thr Ser Ala Leu Leu Gly Ala Arg Leu Lys Glu Leu Glu Gln Glu Ala His Phe Val Ala Gly Glu Arg Phe Leu lle Thr Ser Asn Asn Gln Leu Arg Glu lle Leu Phe Gly Lys Leu Lys Leu His Leu Leu Ser Gln Arg Asn Ser Leu Pro Arg

Thr Gly Leu Gln Lys Tyr Pro Ser Thr Ser Glu Ala Val Asn Ile Gln

Gly Ile Ser Lys His Pro lle Gln Ile Thr Thr Pro Lys Asn Phe Lys Gly Lys Glu Asp Lys lle Leu Thr lle Ser Pro Arg Ala Met Phe Val Ser Ser Lys Gly His Thr Phe Leu Ala Ala Asp Phe Ser Gln Ile Glu Leu Arg Ile Leu Thr His Leu Ser Gly Asp Pro Glu Leu Leu Lys Leu Phe Gln Glu Ser Glu Arg Asp Asp Val Phe Ser Thr Leu Thr Ser Gln Trp Lys Asp Val Pro Val Glu Gln Val Thr His Ala Asp Arg Glu Gln Thr Lys Lys Val Val Tyr Ala Val Val Tyr Gly Ala Gly Lys Glu Arg Leu Ala Ala Cys Leu Gly Val Pro Ile Gln Glu Ala Ala Gln Phe Leu Glu Ser Phe Leu Gln Lys Tyr Lys Lys Ile Lys Asp Phe Ala Arg Ala Ala Ile Ala Gln Cys His Gln Thr Gly Cys Val Val Ser Ile Met Gly Arg Arg Arg Pro Leu Pro Arg Ile His Ala His Asp Gln Gln Leu Arg Ala Gln Ala Glu Arg Gln Ala Val Asn Phe Val Val Gln Ala Gln Ser Gln His Leu Cys Val Glu Val Pro

<210> 4719

<211> 123

<212> PRT

<213> Homo sapiens

<400> 4719

Met Pro Gly Pro Pro Gly Ser Leu Glu Met Gly Pro Leu Thr Phe Arg

1 5 10 15

Asp Val Ala Ile Glu Phe Ser Leu Glu Glu Trp Gln Cys Leu Asp Thr 25 Ala Gln Arg Asn Leu Tyr Arg Lys Val Met Phe Glu Asn Tyr Arg Asn 45 35 40 Leu Val Phe Leu Gly Ile Ala Val Ser Lys Pro His Leu Ile Thr Cys 60 55 Leu Glu Gln Gly Lys Glu Pro Trp Asn Arg Lys Arg Gln Glu Met Val 70 75 Ala Lys Pro Pro Glu Ser Tyr Cys Val Ala Gln Ala Asp Leu Glu Leu 85 Leu Val Ser Ser Tyr Leu Thr Ala Leu Ala Ser Leu Lys Met Trp Asp 105 Tyr Arg Asn Asn Pro Leu Cys Gln Ala Thr Met 120 115

<210> 4720

<211> 301

<212> PRT

<213> Homo sapiens

<400> 4720

Met Ala Val Arg Pro Gly Leu Trp Pro Ala Leu Leu Gly Ile Val Leu

1 5 10 15

Ala Ala Trp Leu Arg Gly Ser Gly Ala Gln Gln Ser Ala Thr Val Ala
20 25 30

Asn Pro Val Pro Gly Ala Asn Pro Asp Leu Leu Pro His Phe Leu Val

35 40 45

Glu Pro Glu Asp Val Tyr Ile Val Lys Asn Lys Pro Val Leu Leu Val 50 55 60

Cys Lys Ala Val Pro Ala Thr Gln Ile Phe Phe Lys Cys Asn Gly Glu
65 70 75 80

Trp Val Arg Gln Val Asp His Val 11e Glu Arg Ser Thr Asp Gly Ser 85 90 95

Asn Gly Leu Pro Thr Met Glu Val Arg Ile Asn Val Ser Arg Gln Gln
100 105 110

Val Glu Lys Val Phe Gly Leu Glu Glu Tyr Trp Cys Gln Cys Val Ala Trp Ser Ser Ser Gly Thr Thr Lys Ser Gln Lys Ala Tyr lle Arg lle Ala Tyr Leu Arg Lys Asn Phe Glu Gln Glu Pro Leu Ala Lys Glu Val Ser Leu Glu Gln Gly Ile Val Leu Pro Cys Arg Pro Pro Glu Gly Ile Pro Pro Ala Glu Val Glu Trp Leu Arg Asn Glu Asp Leu Val Asp Pro Ser Leu Asp Pro Asn Val Tyr Ile Thr Arg Glu His Ser Leu Val Val Arg Gln Ala Arg Leu Ala Asp Thr Ala Asn Tyr Thr Cys Val Ala Łys Asn Ile Val Ala Arg Arg Arg Ser Ala Ser Ala Ala Val lle Val Tyr Val Asp Gly Ser Trp Ser Pro Trp Ser Lys Trp Ser Ala Cys Gly Leu Asp Cys Thr His Trp Arg Ser Arg Glu Cys Ser Asp Pro Ala Pro Arg Asn Gly Gly Glu Glu Cys Gln Gly Thr Asp Leu Asp Thr Arg Asn Cys Thr Ser Asp Leu Cys Val His Ser Glu Ser Ser Leu Pro

<210> 4721

<211> 101

<212> PRT

<213> Homo sapiens

<400> 4721

Met Leu Arg Glu Val Cys Val Cys Val Cys Val Cys Val Cys Val Cys I Cys Val Cys Val Cys I Cy

Pro Glu Ser Trp Arg Phe Thr Phe Cys Lys Gly Arg Gly His Ser Trp 40 Thr Val Thr Ile Pro Ile Leu Val Arg Val Ala Gly Thr Glu Gln Ser 50 55 60 His Gln His Ala Lys Val His Leu Gln Ser Ser Met His Ala Pro Arg 70 75 Lys Pro Pro Val Gly Tyr Ala Ser Cys Thr Phe Pro Phe Ser Leu Thr 90 Ser Val Ser Cys Leu 100 <210> 4722 <211> 116 <212> PRT <213> Homo sapiens <400> 4722 Met Thr Phe Arg Ser Gly Gly Gly Asp Ala Leu Gly Lys Ala Ala Cys 1 5 15 Leu Val Pro Ala Ala Ser Arg Pro Gln Ser Pro Ile Leu Gln Met Thr 25 Ser Arg Glu Val Thr Pro Gln Val Gly Asn Arg Ala Gly Phe Gly Ile 35 40 45 Arg Val Cys Leu Phe Pro Pro Arg Asp Pro Glu Ser Trp Gln Pro Val 55 60 Ser Lys Leu Leu Tyr Val His His Thr Ser Gly Phe Arg Phe Ile Gly 70 75 Val Phe Leu Lys Leu Arg Leu Val Ser Val Gln Leu Leu Leu Val Arg 90 85 95 His Leu Ser Tyr Thr Arg His Cys Pro Trp Cys Trp Arg His Ser Asn 100 105 110

Glu Glu Asp Arg 115

```
⟨210⟩ 4723
<211> 166
<212> PRT
<213> Homo sapiens
⟨400⟩ 4723
Met Arg His Thr Ser Lys Arg Lys Pro Gln Tyr Tyr Glu Ala Glu Met
                  5
                                     10
Val Leu Lys Tyr Tyr Lys His Leu Glu Glu Gly Ser Val Ser Leu Cys
                                                      30
             20
                                 25
Cys Pro Gly Trp Ser Ala Val Ala Gln Ser Arg Leu Thr Ala Ala Ser
Thr Phe Gly Ala Gln Val lle Leu Leu Phe Gln Leu Ser Glu Gln Leu
     50
                         55
                                              60
Arg Leu Gln Glu Leu Lys Leu Pro Thr Phe Arg Ala His Ser Pro Leu
                     70
                                          75
 65
Leu Lys Ser Arg Arg Phe Phe Val Asp Ile Leu Thr Leu Leu Ser Ser
                                     90
                 85
His Cys Gln Leu Cys Pro Ala Ala Arg His Leu Ala Val Tyr Leu Leu
                                 105
                                                     110
            100
Asp His Phe Met Asp Arg Tyr Asn Val Thr Thr Ser Lys Gln Leu Tyr
                            120
Thr Val Ala Val Ser Cys Leu Leu Leu Ala Ser Arg Asn Lys Gly Ser
                        135
                                             140
Gly Ser Pro Val Pro Thr Arg Ser Ala Gln Gln Cys Arg Gln Thr Trp
                                                             160
145
                    150
                                         155
Ala Arg Gly Ser Pro Trp
```

<210> 4724

<211> 165

<212> PRT

<213> Homo sapiens

165

<400> 4724

Met Leu Phe Phe Asn Lys Lys Asn Phe Phe Asn His Gly Leu Ser Gly 10 Phe Ser Cys Pro Leu Asp Thr Phe Leu Cys Leu Ser Leu Ser Leu Phe 20 25 30 Pro Ala Leu His Arg Gly Pro Pro Gly Ser Arg Gly Pro Leu Ile Pro Pro Leu Leu Ser Leu Pro Pro Pro Pro Trp Gly Arg Gly Pro Ile Arg 55 Arg Gly Leu Gly Pro Arg Ser Ser Pro Tyr Gly Arg Gly Trp Trp Gly 75 70 Val Asn Ala Glu Pro Pro Phe Pro Gly Pro Gly His Gly Gly Pro Thr 90 Arg Gly Ser Phe His Lys Glu Gln Arg Asn Pro Arg Arg Leu Lys Ser 100 105 110 Trp Ser Leu lle Lys Asn Thr Cys Pro Pro Lys Asp Pro Gln Val 115 120 125 Met Glu Asp Lys Ser Asp Arg Pro Val Cys Arg His Phe Ala Lys Lys 135 Gly His Cys Arg Tyr Glu Asp Leu Cys Ala Phe Tyr His Pro Gly Val 145 150 155 160 Asn Gly Pro Pro Leu 165

<210> 4725 <211> 141

<212> PRT

<213> Homo sapiens

35

<400> 4725

40

Gly Ser Arg Asp Ala Thr Leu Arg Val Trp Asp Ile Glu Thr Gly Gln Cys Leu His Val Leu Met Gly His Val Ala Ala Val Arg Cys Val Gln Tyr Asp Gly Arg Arg Val Val Ser Gly Ala Tyr Asp Phe Met Val Lys Ala Trp Asp Pro Glu Thr Glu Thr Cys Leu His Thr Leu Gln Gly His Thr Asn Arg Val Tyr Ser Leu Gln Val Arg Ser Leu Ile Ser Pro Leu Asn Ala Leu Leu Met Asn His Lys Val Val Leu Leu Arg

<210> 4726

<211> 326

<212> PRT

<213> Homo sapiens

<400> 4726

Met Asp Ala Ile Lys Lys Lys Met Gln Met Leu Lys Leu Asp Lys Glu Asn Ala Leu Asp Arg Ala Glu Gln Ala Glu Ala Asp Lys Lys Ala Ala Glu Asp Arg Ser Lys Gln Leu Glu Glu Asp lle Ala Ala Lys Glu Lys Leu Leu Arg Val Ser Glu Asp Glu Arg Asp Arg Val Leu Glu Glu Leu His Lys Ala Glu Asp Ser Leu Leu Ala Ala Glu Glu Ala Ala Ala Lys Leu Glu Asp Glu Leu Val Ser Leu Gln Lys Lys Leu Lys Gly Thr Glu Asp Glu Leu Asp Lys Tyr Ser Glu Ala Leu Lys Asp Ala Gln Glu Lys

Leu Glu Leu Ala Glu Lys Lys Ala Thr Asp Ala Glu Ala Asp Val Ala

Ser Leu Asn Arg Arg Ile Gln Leu Val Glu Glu Leu Asp Arg Ala Gln Glu Arg Leu Ala Thr Ala Leu Gln Lys Leu Glu Glu Ala Glu Lys Ala Ala Asp Glu Ser Glu Arg Gly Met Lys Val Ile Glu Ser Arg Ala Gln Lys Asp Glu Glu Lys Met Glu Ile Gln Glu Ile Gln Leu Lys Glu Ala Lys His Ile Ala Glu Asp Ala Asp Arg Lys Tyr Glu Glu Val Ala Arg Lys Leu Val Ile Ile Glu Ser Asp Leu Glu Arg Ala Glu Glu Arg Ala Glu Leu Ser Glu Gly Lys Cys Ala Glu Leu Glu Glu Glu Leu Lys Thr Val Thr Asn Asn Leu Lys Ser Leu Glu Ala Gln Ala Glu Lys Tyr Ser Gln Lys Glu Asp Arg Tyr Glu Glu Glu Ile Lys Val Leu Ser Asp Lys Leu Lys Glu Ala Glu Thr Arg Ala Glu Phe Ala Glu Arg Ser Val Thr Lys Leu Glu Lys Ser Ile Asp Asp Leu Glu Glu Lys Val Ala His Ala Lys Glu Glu Asn Leu Ser Met His Gln Met Leu Asp Gln Thr Leu Leu Glu Leu Asn Asn Met

<210> 4727

<211> 268

<212> PRT

<213> Homo sapiens

<400> 4727

Met Ala Ala Thr Gly Ala Val Ala Ala Ser Ala Ala Ser Gly Gln

1				5					10					15	
Ala	Glu	Gly	Lys	Lys	Ile	Thr	Asp	Leu	Arg	Val	lle	Asp	Leu	Lys	Ser
			20					25					30		
Glu	Leu	Lys	Arg	Arg	Asn	Leu	Asp	11e	Thr	Gly	Val	Lys	Thr	Val	Leu
		35					40					45			
lle	Ser	Arg	Leu	Lys	Gln	Ala	lle	Glu	Glu	Glu	Gly	Gly	Asp	Pro	Asp
	50					55					60				
Asn	lle	Glu	Leu	Thr	Val	Ser	Thr	Gly	Thr	Pro	Asn	Lys	Lys	Pro	Thr
65					70					75					80
Lys	Gly	Lys	Gly	Lys	Lys	His	Glu	Ala	Asp	Glu	Leu	Ser	Gly	Asp	Ala
				85					90					95	
Ser	Val	Glu	Asp	Asp	Ala	Phe	He	Lys	Asp	Gly	Glu	Glu	Glu	Glu	Asn
			100					105					110		
Glu	Lys	Gly	Ser	Leu	Ala	Glu	Ala	Asp	His	Thr	Ala	His	Glu	Glu	Met
		115					120					125			
Glu	Ala	His	Thr	Thr	Val	Lys	Glu	Ala	Glu	Asp	Asp	Asn	lle	Ser	Val
	130					135					140				
Thr	Ile	Gln	Ala	Glu	Asp	Ala	lle	Thr	Leu	Asp	Phe	Asp	Gly	Asp	Asp
145					150					155					160
Leu	Leu	Glu	Thr	Gly	Lys	Asn	Val	Lys	lle	Thr	Asp	Ser	Glu	Ala	Ser
				165					170					175	
Lys	Pro	Lys	Asp	Gly	Gln	Asp	Ala	He	Ala	Gln	Ser	Pro	Glu	Lys	G] u
			180					185					190		
Ser	Lys	Asp	Tyr	Glu	Met	Asn	Ala	Asn	His	Lys	Asp	Gly	Lys	Lys	Glu
		195					200					205			
Asp	Cys	Val	Lys	Gly	Asp	Pro	Val	Glu	Lys	Glu	Ala	Arg	Glu	Ser	Ser
	210					215					220				
Lys	Lys	Ala	Glu	Ser	Gly	Asp	Lys	Glu	Lys	Asp	Thr	Leu	Lys	Lys	Gly
225					230					235					240
Pro	Ser	Ser	Thr	Gly	Ala	Ser	Gly	Gln	Ala	Lys	Arg	Phe	Val	Phe	Leu
				245					250					255	
Cys	Gln	Phe	Phe	Thr	He	Leu	Asn	Ser	Ser	11e	Gln				
			260					265							

```
<210> 4728
<211> 105
<212> PRT
<213> Homo sapiens
<400> 4728
Met Thr Thr His Leu Tyr Asp Ala Pro Thr Val Lys Phe Leu Thr Pro
 1
                  5
                                     10
Cys Tyr His Pro Asn Val Asp Thr Gln Gly Asn Ile Cys Leu Asp Ile
             20
                                 25
                                                      30
Leu Lys Glu Lys Trp Ser Ala Pro Tyr Asp Ile Arg Thr Ile Leu Leu
                             40
Ser 11e Gln Cys Leu Leu Gly Gln Leu Asn 11e Asp Ser Pro Leu Asn
     50
                         55
                                              60
Thr His Ala Thr Lys Leu Trp Glu Asn Pro Ile Ala Leu Arg Ser Thr
65
                     70
                                          75
Cys Lys Gly Gln Ala Gln Trp Leu Thr Pro Val Ile Pro Ala Leu Trp
                 85
                                     90
                                                          95
Glu Ala Glu Val Gly Gly Trp Ile Thr
            100
                                 105
<210> 4729
⟨211⟩ 336
<212> PRT
<213> Homo sapiens
<400> 4729
Met Lys Glu Arg Lys Arg His Leu Gly Asp Thr Lys His Phe Cys Pro
1
                  5
                                      10
                                                          15
Val Val Leu Lys Glu Asn Phe lle Leu Gln Pro Gly Asn Thr Glu Glu
                                 25
Ala Ala Lys Tyr Arg Glu Lys Ile Tyr Tyr Phe Ser Ser Ala Glu Ala
         35
                                                  45
                             40
Lys Glu Lys Phe Leu Glu His Pro Glu Asp Tyr Val Ala His Glu Glu
```

55

60

Pro	Leu	Lys	Ala	Pro	Pro	Leu	Arg	Пe	Cys	Leu	Val	Gly	Pro	Gln	Gly
65					70					75					80
Ser	Gly	Lys	Thr	Met	Cys	Gly	Arg	Gln	Leu	Ala	Glu	Lys	Leu	Asn	lle
				85					90					95	
Phe	His	He	Gln	Phe	Glu	Glu	Val	Leu	Gln	Glu	Lys	Leu	Leu	Leu	Lys
			100					105					110		
Thr	Glu	Lys	Lys	Val	Gly	Pro	Glu	Phe	Glu	Glu	Asp	Ser	Glu	Asn	Glu
		115					120					125			
Gln	Ala	Ala	Lys	Gln	Glu	Leu	Glu	Glu	Leu	Ala	Ile	Gln	Ala	Asn	Val
	130					135					140				
Lys	Val	Glu	Glu	Glu	Asn	Thr	Lys	Lys	G1n	Leu	Pro	Glu	Val	Gln	Leu
145					150					155					160
Thr	Glu	Glu	Glu	Glu	Val	lle	Lys	Ser	Ser	Leu	Met	Glu	Asn	Glu	Pro
				165					170					175	
Leu	Pro	Pro	G1u	He	Leu	Glu	Val	lle	Leu	Ser	Glu	Trp	Trp	Leu	Lys
			180					185					190		
Glu	Pro	He	Arg	Ser	Thr	Gly	Phe	Ile	Leu	Asp	Gly	Phe	Pro	Arg	Tyr
		195					200					205			
Pro	Glu	Glu	Ala	Gln	Phe	Leu	Gly	Asp	Arg	Gly	Phe	Phe	Pro	Asp	Ala
	210					215					220				
Ala	Val	Phe	lle	Gln	Val	Asp	Asp	Gln	Asp	He	Phe	Asp	Arg	Leu	
225					230					235					240
Pro	Ala	Gln	lle		Lys	Trp	Lys	Leu		Gln	Lys	Lys	Lys	Leu	Glu
				245					250					255	
Arg	Lys	Lys		He	Lys	Asp	Met		Ala	Lys	He	Arg		Asp	Thr
			260					265					270		
He	Ala		Arg	Arg	Ala	Glu		lle	Leu	Glu	Arg		Lys	Lys	Arg
		275	_				280					285	_		_
Arg		Val	Ser	Ser	Phe		Phe	Phe	Phe	Lys		Gly	Ser	His	Ser
	290					295	_				300				
	Ala	Gln	Gly	Arg		GIn	Trp	His	Asn		Ser	Ser	Leu	Gln	
305			۵.		310					315	~				320
Arg	Thr	Pro	Gly		Lys	Gly	Ser	Ser		Leu	Ser	Leu	Ser	Lys	Cys
				325					330					335	

```
<210> 4730
<211> 438
<212> PRT
<213> Homo sapiens
<400> 4730
Met His Pro Leu Pro Gly Tyr Trp Ser Cys Tyr Cys Leu Leu Leu Leu
                  5
                                     10
Phe Ser Leu Gly Val Gln Gly Ser Leu Gly Ala Pro Ser Ala Ala Pro
             20
                                 25
                                                      30
Glu Gln Val His Leu Ser Tyr Pro Gly Glu Pro Gly Ser Met Thr Val
Thr Trp Thr Trp Val Pro Thr Arg Ser Glu Val Gln Phe Gly Leu
     50
                         55
                                              60
Gln Pro Ser Gly Pro Leu Pro Leu Arg Ala Gln Gly Thr Phe Val Pro
                     70
                                          75
65
Phe Val Asp Gly Gly Ile Leu Arg Arg Lys Leu Tyr Ile His Arg Val
                                     90
                 85
Thr Leu Arg Lys Leu Leu Pro Gly Val Gln Tyr Val Tyr Arg Cys Gly
            100
                                 105
                                                     110
Ser Ala Gln Gly Trp Ser Arg Arg Phe Arg Phe Arg Ala Leu Lys Asn
                            120
Gly Ala His Trp Ser Pro Arg Leu Ala Val Phe Gly Asp Leu Gly Ala
                                             140
                        135
Asp Asn Pro Lys Ala Val Pro Arg Leu Arg Arg Asp Thr Gln Gln Gly
                    150
                                         155
                                                             160
145
Met Tyr Asp Ala Val Leu His Val Gly Asp Phe Ala Tyr Asn Leu Asp
                165
                                     170
Gln Asp Asn Ala Arg Val Gly Asp Arg Phe Met Arg Leu Ile Glu Pro
                                                     190
            180
                                 185
Val Ala Ala Ser Leu Pro Tyr Met Thr Cys Pro Gly Asn His Glu Glu
                            200
                                                 205
Arg Tyr Asn Phe Ser Asn Tyr Lys Ala Arg Phe Ser Met Pro Gly Asp
    210
                        215
                                             220
Asn Glu Gly Leu Trp Tyr Ser Trp Asp Leu Gly Pro Ala His Ile Ile
```

Ser Phe Ser Thr Glu Val Tyr Phe Phe Leu His Tyr Gly Arg His Leu Val Gln Arg Gln Phe Arg Trp Leu Glu Ser Asp Leu Gln Lys Ala Asn Lys Asn Arg Ala Ala Arg Pro Trp Ile Ile Thr Met Gly His Arg Pro Met Tyr Cys Ser Asn Ala Asp Leu Asp Asp Cys Thr Arg His Glu Ser Lys Val Arg Lys Gly Leu Gln Gly Lys Leu Tyr Gly Leu Glu Asp Leu Phe Tyr Lys Tyr Gly Val Asp Leu Gln Leu Trp Ala His Glu His Ser Tyr Glu Arg Leu Trp Pro lle Tyr Asn Tyr Gln Val Phe Asn Gly Ser Arg Glu Met Pro Tyr Thr Asn Pro Arg Gly Pro Val His Ile Ile Thr Gly Ser Ala Gly Cys Glu Glu Arg Leu Thr Pro Phe Ala Val Phe Pro Arg Pro Trp Ser Ala Val Arg Val Lys Glu Tyr Gly Tyr Thr Arg Leu His lle Leu Asn Gly Thr His Thr His lle Gln Gln Val Ser Asp Asp Gln Asp Gly Lys Ile Val Asp Asp Val Trp Val Val Arg Pro Leu Phe Gly Arg Arg Met Tyr Leu

<210> 4731

<211> 607

<212> PRT

<213> Homo sapiens

<400> 4731

Met Ala Glu Glu Gly Ala IIe Leu Lys Val Thr Lys Asp Leu Arg Ala

1				5					10					15	
Ala	Val	Ser	Ala	He	Leu	Gln	Gly	Tyr	G1 y	Asp	Gly	Gln	Gly	Pro	Val
			20					25					30		
Thr	Asp	Thr	Ser	Ala	Glu	Leu	His	Arg	Leu	Cys	Gly	Cys	Leu	Glu	Leu
		35					40					45			
Leu	Leu	Gln	Phe	Asp	Gln	Lys	Glu	Gln	Lys	Ser	Phe	Leu	Gly	Pro	Arg
	50					55					60				
Lys	Asp	Tyr	Trp	Asp	Phe	Leu	Cys	Thr	Ala	Leu	Λrg	Arg	Gln	Arg	G] y
65					70					75					80
Asn	Met	Glu	Pro	Ile	His	Phe	Val	Arg	Ser	Gln	Asp	Lys	Leu	Lys	Thr
				85					90					95	
Pro	Leu	Gly	Lys	Gly	Arg	Ala	Phe	He	Arg	Phe	Cys	Leu	Ala	Arg	Gly
			100					105					110		
Gln	Leu	Ala	Glu	Ala	Leu	Gln	Leu	Cys	Leu	Leu	Asn	Ser	Glu	Leu	Thr
		115					120					125			
Arg	Glu	Trp	Tyr	Gly	Pro	Arg	Ser	Pro	Leu	Leu	Cys	Pro	Glu	Arg	Gln
	130					135					140				
Glu	Asp	Ile	Leu	Asp	Ser	Leu	Tyr	Ala	Leu	Asn	Gly	Val	Ala	Phe	Glu
145					150					155					160
Leu	Asp	Leu	Gln	Gln	Pro	Asp	Leu	Asp	Gly	Ala	Trp	Pro	Met	Phe	Ser
				165					170					175	
Glu	Ser	Arg	Cys	Ser	Ser	Ser	Thr	Gln	Thr	Gln	Gly	Arg	Arg	Pro	Arg
			180					185					190		
Lys	Asn	Lys	Asp	Ala	Pro	Lys	Lys	Val	Pro	Leu	Pro	Cys	Leu	His	Ser
		195					200					205			
Glu	Leu	Pro	Leu	Pro	His	Leu	Leu	Gly	He	Leu	Lys	He	Pro	Ala	Ala
	210					215					220				
Tyr	Gly	Gly	Pro	Glu	Asn	Val	Gln	11e	Glu	Asp	Ser	llis	Thr	Ser	Gln
225					230					235					240
Ala	lle	Cys	Leu	Gln	Asp	Ala	Pro	Ser	Gly	Gln	Gln	Leu	Ala	G] y	Leu
				245					250					255	
Pro	Arg	Ser	Gln	GIn	Gln	Arg	His	Leu	Pro	Phe	Phe	Leu	Glu	Lys	Lys
			260					265					270		
G1 y	Glu	Ser	Ser	Arg	Lys	His	Arg	Tyr	Pro	GIn	Ser	Met	Trp	Glu	Pro
		275					280					285			
Glu	Gly	Lys	Glu	Leu	GIn	l.eu	Asp	Gln	$Gl\mathbf{u}$	Glu	Arg	Ala	Pro	Trp	He

	290					295					300				
Glu	lle	Phe	Leu	G1 y	Asn	Ser	Thr	Pro	Ser	Thr	Gln	G] y	Gln	Gly	Lys
305					310					315					320
Gly	Ala	Met	Gly	Thr	Gln	Lys	Glu	Val	11e	Gly	Met	G] u	Ala	Glu	Val
				325					330					335	
Thr	Gly	Val	Leu	Leu	Val	Ala	Glu	Gly	Gln	Arg	Thr	Thr	Glu	Gly	Thr
			340					345					350		
His	Lys	Lys	Glu	Ala	Glu	Trp	Ser	His	Val	Gln	Arg	Leu	Leu	Met	Pro
		355					360					365			
Ser	Pro	Arg	Gly	Ala	Val	Glu	G1 y	Ala	Val	Ser	Gly	Ser	Arg	Gln	Gly
	370					375					380				
Ser	Gly	Gly	Ser	Ser	lle	Leu	Gly	Glu	Pro	Trp	Val	Leu	Gln	Gly	His
385					390					395					400
Ala	Thr	Lys	Glu	Asp	Ser	Thr	Val	Glu	Asn	Pro	Gln	Val	Gln	Thr	Glu
				405					410					415	
Val	Thr	Leu	Val	Ala	Arg	Arg	Glu	Glu	Gln	Ala	Glu	Val	Ser	Leu	Gln
			420					425					430		
Asp	Glu	Ile	Lys	Ser	Leu	Arg	Leu	Gly	Leu	Arg	Lys	Ala	Glu	Glu	Gln
		435					440					445			
Λla	Gln	Arg	Gln	Glu	Gln	Leu	Leu	Arg	Glu	Gln	Glu	Gly	Glu	Leu	Gln
	450					455					460				
Ala	Leu	Arg	Glu	Gln	Leu	Ser	Arg	Cys	Gln	Glu	Glu	Arg	Ala	Glu	Leu
465					470					475					480
Gln	Ala	G1n	Leu	Glu	Gln	Lys	Gln	Gln	Glu	Ala	Glu	Arg	Arg	Asp	Ala
				485					490					495	
Met	Tyr	Gln	Glu	Glu	Leu	Gly	Gly	Gln	Arg	Asp	Leu	Val	Gln	Ala	Met
			500					505					510		
Lys	Arg		Val	Leu	Glu	Leu		Gln	Glu	Lys	Asp		Leu	Trp	Gln
		515					520					525			
Arg		Gln	His	Leu	Ser		Met	Ala	Pro	Glu		Cys	Vał	Ala	Cys
	530					535					540				
	Lys	He	Phe	Gly		Phe	Ser	Arg	Arg		Pro	Cys	Ser	Ala	
545					550					555					560
Thr	Ser	Leu	Gly		Glu	Glu	Ser	Glu		Pro	Gly	G1 u	Ala	Ala	Met
T		T.		565	T		63		570	C	0.1		15	575	C.1
Trp	Arg	Trp	Arg	Leu	Irp	Arg	Gly	Arg	Ala	Ser	GLy	His	Pro	Ser	GLy

| S80 | S85 | S90 | S90 | Glu Gly Arg Pro Pro Glu Thr Asn Gln Gly His Ser Thr Gly Arg | S95 | 600 | 605 |

<210> 4732

<211> 148

<212> PRT

<213> Homo sapiens

<400> 4732

Met Ser Cys Asn Leu Ala Ser Leu Cys Ser Lys Val Leu Glu Gln Gly
1 5 10 15

Ala Tyr His Pro 11e Leu 11e Pro Ser Pro Leu Ser Pro Pro Trp Thr
20 25 30

His Cys Ala Gln His Leu Arg Pro Cys Thr Cys Lys Gly Ser Ser His
35 40 45

Tyr Gly Asp Gln Gln Pro Arg Ser Gln Val Leu Asn Ser Lys Met Leu 50 55 60 .

Leu Ser Ala Ser Val Leu Asp Leu Ser Ser Val Asp Ser Asp Ala Phe
65 70 75 80

Glu Ile Leu Leu Ser Leu Ala Ser Ile Ala Ile Arg Thr Pro Arg Val 85 90 95

Cys His Pro Leu Leu Pro Leu His Phe 11e Leu Leu His Lys Gly Gln 100 105 110

Pro Ser His Val Glu Val Leu Trp Asp Ala Ala Trp Leu Phe Phe 11e 115 120 125

Val Asp Leu Met Phe Ser Ser Ile Leu Leu Cys Thr Phe Leu Arg Leu 130 135 140

Gln Tyr Phe Val

145

<210> 4733

<211> 392

<212> PRT

<213> Homo sapiens

<400)> 47	733													
Met	Gln	Pro	Leu	Thr	Lys	Asp	Ala	Gly	Met	Ser	Leu	Ser	Ser	Val	Thr
1				5					10					15	
Leu	Ala	Ser	Ala	Leu	Gln	Val	Arg	Gly	Glu	Ala	Leu	Ser	Glu	Glu	Glu
			20					25					30		
Ile	Trp	Ser	Leu	Leu	Phe	Leu	Ala	Ala	Glu	Gln	Leu	Leu	Glu	Asp	Leu
		35					40					45			
Arg	Asn	Asp	Ser	Ser	Asp	Tyr	Val	Va]	Cys	Pro	Trp	Ser	Ala	Leu	Leu
	50					55					60				
Ser	Ala	Ala	Gly	Ser	Leu	Ser	Phe	Gln	Gly	Arg	Val	Ser	His	Ile	Glu
65					70					75					80
Ala	Ala	Pro	Phe	Lys	Ala	Pro	Glu	Leu	Leu	Gln	Gly	Gln	Ser	Glu	Asp
				85					90					95	
Glu	Gln	Pro	Asp	Ala	Ser	Gln	Met	His	Va]	Tyr	Ser	Leu	Gly	Met	Thr
			100					105					110		
Leu	Tyr	Trp	Ser	Ala	Gly	Phe	His	Val	Pro	Pro	His	Gln	Pro	Leu	Gln
		115					120					125			
Leu	Cys	Glu	Pro	Leu	His	Ser	He	Leu	Leu	Thr	Met	Cys	Glu	Asp	Gln
	130					135					140				
Pro	His	Arg	Arg	Cys	Thr	Leu	Gln	Ser	Val	Leu	Glu	Ala	Cys	Arg	Val
145					150					155					160
His	Glu	Lys	Glu	Val	Ser	Val	Tyr	Pro	Ala	Pro	Ala	Gly	Leu	His	He
				165					170					175	
Arg	Arg	Leu		Gly	Leu	Val	Leu		Thr	lle	Ser	Glu		Glu	Lys
			180					185					190		
Arg	Val		Glu	Glu	Ser	Ser		Val	Gln	Gln	Asn		Ser	Tyr	Leu
		195					200		_		_	205			
Leu		Lys	Arg	Leu	Arg		Thr	Ser	Ser	61u		Pro	Ala	Ala	Gln
	210					215	_				220		_		
	Pro	Glu	Cys	Leu		Pro	Cys	Arg	Val		Glu	Arg	Ser	Thr	
225		_			230			_		235					240
Thr	GIn	Ser	Ser		Glu	Pro	His	Trp		Thr	Leu	Thr	His	Ser	His
0	0			245					250	61			T>.	255	
Cys	Ser	Leu	Leu	Val	Asn	Arg	Ala	Leu	Pro	ыу	Ala	Asp	Pro	Gln	Asp

	260			265					270		
Gln Gln Ala	Gly Arg	Arg Leu	Ser	Ser	Gly	Ser	Val	His	Ser	Ala	Ala
275			280					285			
Asp Ser Ser	Trp Pro	Thr Thr	Pro	Ser	Gln	Arg	Gly	Phe	Leu	Gln	Arg
290		295					300				
Arg Ser Lys	Phe Ser	Arg Pro	Glu	Phe	Пe	Leu	Leu	Ala	Gly	Glu	Ala
305		310				315					320
Pro Met Thr	Leu His	Leu Pro	Gly	Ser	Val	Val	Thr	Lys	Lys	Gly	Lys
	325				330					335	
Ser Tyr Leu	Ala Leu	Arg Asp	Leu	Cys	Val	Val	Leu	Leu	Asn	Gly	Gln
	340			345					350		
His Leu Glu	Val Lys	Cys Asp	Va]	Glu	Ser	Thr	Val	Gly	Ala	Val	Phe
355			360					365			
Asn Ala Val	Thr Ser	Phe Ala	Asn	Leu	Glu	Glu	Leu	Thr	Tyr	Phe	Gly
370		375					380				
Leu Ala Tyr	Met Lys	Ser Gly	Glu								
385		390									

<210> 4734

<211> 357

<212> PRT

<213> Homo sapiens

<400> 4734

Met Pro Glu Cys Trp Asp Gly Glu His Asp Ile Glu Thr Pro Tyr Gly 10 Leu Leu His Val Val Ile Arg Gly Ser Pro Lys Gly Asn Arg Pro Ala 20 25 30 lle Leu Thr Tyr His Asp Val Gly Leu Asn Arg Lys Cys Ser Pro Ala 35 45 Ser Val Ser Pro Pro Leu Pro Pro Ile Ser Gln Ser Asp Lys Leu Cys 55 Phe Asn Thr Phe Phe Asn Phe Glu Asp Met Gln Glu lle Thr Lys His 75 65 70 80

Phe Val Val Cys His Val Asp Ala Pro Gly Gln Gln Val Gly Ala Ser

				85					90					95	
Gln	Phe	Pro	Gln	Gly	Tyr	Gln	Phe	Pro	Ser	Met	Glu	Gln	Leu	Ala	Ala
			100					105					110		
Met	Leu	Pro	Ser	Val	Val	Gln	His	Phe	Gly	Phe	Lys	Tyr	Val	He	Gly
		115					120					125			
Ile	Gly	Val	Gly	Ala	Gly	Ala	Tyr	Val	Leu	Ala	Lys	Phe	Ala	Leu	lle
	130					135					140				
Phe	Pro	Asp	Leu	Val	Glu	Gly	Leu	Val	Leu	Val	Asn	He	Asp	Pro	Asn
145					150					155					160
Gly	Lys	Gly	Trp	Ile	Asp	Trp	Ala	Ala		Lys	Leu	Ser	Gly		Thr
				165					170					175	
Ser	Thr	Leu		Asp	Thr	Val	Leu		His	Leu	Phe	Ser	Gln	Glu	Glu
			180					185				0.1	190		61
Leu	Val		Asn	Thr	Glu	Leu		GIn	Ser	Tyr	Arg		Gln	11e	Gly
	17.1	195		C1.	۸3.	Α	200	C1	1	DI	т	205	М.,	Т	Α
Asn		vai	Asn	GIn	Ala		Leu	GIN	Leu	Pne		Asn	Met	ıyr	Asn
Con	210	Ana	A on	Lou	A an	215	Aan	Ana	Dro	C1.	220 Thr	Vo.1	Dro	Acn	110
225	nı g	лід	nsh	Leu	230	116	USII	A1 g	110	235	1111	vai	Pro	nsii	240
	Thr	Len	Arø	Cvs		Val	Met	Leu	Val		Glv	Asp	Asn	Ala	
12,5		Bea	111 8	245	110	, , ,	Mec	Leu	250	741	0.19	пор	71071	255	
Ala	Glu	Asp	Glv		Val	Glu	Cvs	Asn		Lvs	Leu	Asp	Pro		Thr
			260				,	265		Ž		•	270		
Thr	Thr	Phe	Leu	Lys	Met	Ala	Asp	Ser	Gly	Gly	Leu	Pro	Gln	Val	Thr
		275					280					285			
Gln	Pro	Gly	Lys	Leu	Thr	Glu	Ala	Phe	Lys	Tyr	Phe	Leu	Gln	Gly	Met
	290					295					300				
Gly	Tyr	Met	Pro	Ser	Ala	Ser	Met	Thr	Arg	Leu	Ala	Arg	Ser	Arg	Thr
305					310					315					320
Ala	Ser	Leu	Thr	Ser	Ala	Ser	Ser	Val	Asp	Gly	Ser	Arg	Pro	G1n	Ala
				325					330					335	
Cys	Thr	His	Ser	Glu	Ser	Ser	Glu	Gly	Leu	G1 y	Gln	Val	Asn	His	Thr
			340					345					350		
Met	Glu	Val	Ser	Cys											

<210> 4735 <211> 357 <212> PRT <213> Homo sapiens <400> 4735 Met Ser Pro Gly Pro Thr Asn Phe Phe Val Phe Leu Leu Glu Met Gly Phe His His Ala Gly Arg Val Gly Pro Glu Leu Leu Thr Ser Gly Asp Pro Leu Ala Ser Ala Ser Gln Ser Ala Gly Ile Thr Glu Thr Glu Gly Ser Glu Glu Asp Asp Lys Glu Asn Asp Lys Thr Glu Glu Met Pro Asn Asp Ser Val Leu Glu Asn Lys Ser Leu Gln Glu Asn Glu Glu Glu Glu lle Gly Asn Leu Glu Leu Ala Trp Asp Met Pro Asp Leu Ala Lys Ile Ile Phe Lys Arg Gln Glu Thr Lys Glu Ala Gln Leu Tyr Ala Ala Gln Ala His Leu Lys Leu Gly Glu Val Ser Val Glu Ser Glu Asn Tyr Val Gln Ala Val Glu Glu Phe Gln Ser Cys Leu Asn Leu Gln Glu Gln Tyr Leu Glu Ala His Asp Arg Leu Leu Ala Glu Thr His Tyr Gln Leu Gly Leu Ala Tyr Gly Tyr Asn Ser Gln Tyr Asp Glu Ala Val Ala Gln Phe Ser Lys Ser Ile Glu Val lle Glu Asn Arg Met Ala Val Leu Asn Glu Gln Val Lys Glu Ala Glu Gly Ser Ser Ala Glu Tyr Lys Lys Glu Ile Glu Glu Leu Lys Glu Leu Leu Pro Glu 11e Arg Glu Lys 11e Glu Asp

Ala Lys Glu Ser Gln Arg Ser Gly Asn Val Ala Glu Leu Ala Leu Lys 225 230 235 Ala Thr Leu Val Glu Ser Ser Thr Ser Gly Phe Thr Pro Gly Gly Gly 250 255 245 Gly Ser Ser Val Ser Met Ile Ala Ser Arg Lys Pro Thr Asp Gly Ala 260 265 Ser Ser Ser Asn Cys Val Thr Asp Ile Ser His Leu Val Arg Lys Lys 280 275 285 Arg Lys Pro Glu Glu Glu Ser Pro Arg Lys Asp Asp Ala Lys Lys Ala 295 290 Lys Gln Glu Pro Glu Val Asn Gly Gly Ser Gly Asp Ala Val Pro Ser 310 315 Gly Asn Glu Val Ser Glu Asn Met Glu Glu Glu Ala Glu Asn Gln Ala 330 335 325 Glu Ser Arg Ala Ala Val Glu Gly Thr Val Glu Ala Gly Ala Thr Val 340 345 350 Glu Ser Thr Ala Cys 355

<210> 4736

<211> 217

<212> PRT

<213> Homo sapiens

<400> 4736

Met Leu Ser Leu Gln Asp Ser Val Phe Phe Glu 11e Ser 11e Lys Ser 1 5 10 15

Leu Leu Lys Ser Trp Ser Ser Ser Ser Ser Ala Pro Val Ser Lys Val
20 25 30

Asn Lys Tyr Cys Ala Ser Ser Asn Phe His Ser Thr Trp Gly Lys Lys
35 40 45

Asn lle lle Met Ser Asn lle Thr lle Asp Pro Asp Val Lys Pro Gly
50 55 60

Glu Tyr Val Ile Lys Ser Leu Phe Ala Glu Phe Ala Val Gln Ala Glu 65 70 75 80 Lys Lys Ile Glu Val Val Met Ala Glu Pro Leu Glu Lys Leu Leu Ser 90 Arg Ser Leu Gln Arg Gly Glu Asp Leu Gln Phe Asp Gln Leu Ile Ser 100 105 110 Ser Met Ser Ser Val Ala Glu His Cys Leu Pro Ser Leu Leu Arg Thr 120 Leu Phe Asp Trp Tyr Arg Arg Gln Asn Gly Thr Glu Asp Glu Ser Tyr 135 140 Glu Tyr Arg Pro Arg Ser Ser Thr Lys Ser Lys Gly Asp Glu Gln Gln 150 155 160 145 Arg Glu Arg Asp Tyr Leu Leu Glu Arg Arg Asp Leu Ala Val Asp Phe 170 165 Ile Phe Cys Leu Val Leu Val Glu Val Leu Lys Gln Val Ser Ser Phe 185 190 lle Ser Ala Lys Ser Val Phe Leu Ile Leu Cys Leu Leu Cys Leu Arg 195 200 205 Val Ser Val Thr Leu Arg Cys Cys Val 210 215

<210> 4737

<211> 908

<212> PRT

<213> Homo sapiens

<400> 4737

Met Gly Val Asn Asp Leu Trp Gln 11e Leu Glu Pro Val Lys Gln His 1 5 10 15

lle Pro Leu Arg Asn Leu Gly Gly Lys Thr Ile Ala Val Asp Leu Ser 20 25 30

Leu Trp Val Cys Glu Ala Gln Thr Val Lys Lys Met Met Gly Ser Val
35 40 45

Met Lys Pro His Leu Arg Asn Leu Phe Phe Arg Ile Ser Tyr Leu Thr
50 55 60

Gln Met Asp Val Lys Leu Val Phe Val Met Glu Gly Glu Pro Pro Lys 65 70 75 80

Leu	Lys	Ala	Asp	Val	He	Ser	Lys	Arg	Asn	Gln	Thr	Arg	Tyr	Gly	Ser
				85					90					95	
Ser	Gly	Lys	Ser	Trp	Ser	Gln	Lys	Thr	Gly	Arg	Ser	His	Phe	Lys	Ser
			100					105					110		
Val	Leu	Arg	Glu	Cys	Leu	His	Met	Leu	Glu	Cys	Leu	Gly	He	Pro	Trp
		115					120					125			
Val	Gln	Ala	Ala	Gly	Glu	Ala	Glu	Ala	Met	Cys	Ala	Tyr	Leu	Asn	Ala
	130					135					140				
Gly	Gly	His	Val	Asp	Gly	Cys	Leu	Thr	Asn	Asp	Gly	Asp	Thr	Phe	Leu
145					150					155					160
Tyr	Gly	Ala	Gln	Thr	Val	Tyr	Arg	Asn	Phe	Thr	Met	Asn	Thr	Lys	Asp
				165					170					175	
Pro	His	Val	Asp	Cys	Tyr	Thr	Met	Ser	Ser	He	Lys	Ser	Lys	Leu	Gly
			180					185					190		
Leu	Asp	Arg	Asp	Ala	Leu	Va]	Gly	Leu	Ala	Ile	Leu	Leu	Gly	Cys	Asp
		195					200					205			
Tyr	Leu	Pro	Lys	Gly	Val	Pro	Gly	Val	Gly	Lys	Glu	Gln	Ala	Leu	Lys
	210					215					220				
Leu	Ile	Gln	Ile	Leu	Lys	Gly	Gln	Ser	Leu	Leu	Gln	Arg	Phe	Asn	Arg
225					230					235					240
Trp	Asn	Glu	Thr	Ser	Cys	Asn	Ser	Ser	Pro	Gln	Leu	Leu	Val	Thr	Lys
				245					250					255	
Lys	Leu	Ala	His	Cys	Ser	Va]	Cys	Ser	His	Pro	Gly	Ser	Pro	Lys	Asp
			260					265					270		
His	Glu	Arg	Asn	Gly	Cys	Arg	Leu	Cys	Lys	Ser	Asp	Lys	Tyr	Cys	Glu
		275					280					285			
Pro	His	Asp	Tyr	Glu	Tyr	Cys	Cys	Pro	Cys	Glu	Trp	His	Arg	Thr	Glu
	290					295					300				
His	Asp	Arg	Gln	Leu	Asn	Glu	Val	Glu	Asn	Asn	lle	Lys	Lys	Lys	Ala
305					310					315					320
Cys	Cys	Cys	Glu	Gly	Phe	Pro	Phe	His	Glu	Val	He	Gln	Glu	Phe	Leu
				325					330					335	
Leu	Asn	Lys	Asp	Lys	Leu	Val	Lys		He	Arg	Tyr	Gln	Arg	Pro	Asp
			340					345					350		
Leu	Leu		Phe	Gln	Arg	Phe		Leu	Glu	Lys	Met		Trp	Pro	Asn
		355					360					365			

Hi	s T	`yr	Ala	Cys	Glu	Lys	Leu	Leu	Val	Leu	Leu	Thr	His	Tyr	Asp	Met
	3	370					375					380				
11	e G	lu	Arg	Lys	Leu	Gly	Ser	Arg	Asn	Ser	Asn	G1n	Leu	Gln	Pro	Ile
38	5					390					395					400
Ar	g I	le	Val	Lys	Thr	Arg	He	Arg	Asn	G1 y	Val	His	Cys	Phe	Glu	He
					405					410					415	
Gl	u T	`rp	Glu	Lys	Pro	Glu	His	Tyr	Ala	Met	Glu	Asp	Lys	Gln	His	Gly
				420					425					430		
Gl	u P	he		Leu	Leu	Thr	He	Glu	Glu	Glu	Ser	Leu	Phe	Glu	Ala	Ala
			435					440					445			
Тy	r P	ro	Glu	Ile	Val	Ala	Val	Tyr	Gln	Lys	Gln	Lys	Leu	Glu	He	Lys
		150					455					460				
G1	y L	ys	Lys	Gln	Lys	Arg	He	Lys	Pro	Lys	Glu	Asn	Asn	Leu	Pro	Glu
46	5					470					475					480
Pr	o A	sp	Glu	Val		Ser	Phe	Gln	Ser		Met	Thr	Leu	Lys		Thr
					485					490					495	
Су	s G	ilu	He		His	Lys	Gln	Asn		Lys	Leu	Asn	Ser		lle	Ser
				500					505			. 0		510		
Pr	o A	lsp		Thr	Leu	Pro	Gln		Ser	He	Ser	Ala		Leu	Asn	Ser
			515					520	_	_			525			
Le			Leu	Pro	Lys	Asn	Thr	Pro	Cys	Leu	Asn		GIn	Glu	GIn	Phe
		530					535		* 1		0.7	540				
		er	Ser	Leu	Arg		Leu	Ala	He	GIn		He	Lys	Ala	Val	
54				7.7	C	550	c	c	C1	D	555	T)	C	c	11.	560
Ly	s S	er	Leu	11e		Glu	Ser	Ser	GIn		Asn	inr	Ser	Ser		Asn
7.1		,	12 7	7.1	565		,	11.	1	570	TI.	7.1		т	575	C1
11	e S	er	vai		Ala	Asp	Leu	nis		ser	ınr	116	ASP		GIU	GIY
TL	C	`	Dlag	580	A	Can	Dave	A1.c	585	Cln	Ama	1	Than	590	Con	u; o
ın	rs	er		ser	ASN	Ser	Pro	600	116	oin	Arg	ASI		rne	ser	nis
٨٥	ո I	011	595	Son	Clu	Vol	C1		Cl ₁₁	Lou	Sor	Λlα	605	Dro	Aan	Cly
AS			Lys	361	GIU	vai	Glu	261	Olu	Leu	261	620	116	110	игр	ОГУ
D۱		510	Acr	Tla	Dro	C1	615 Gln	Low	Sor	Cvc	Glu		Gl.	Δπα	Tur	The
62		ររូប	lien	лте	110	630	0.111	Leu	261	CyS	635	261	Olu	vi g	1 9 1	640
		len	116	lvc	lvo		Leu	Acr	Glas	Acr		Acr	Glv	110	Sor	
/ 1.1	a P	1911	116	LYS	645	val	Leu	ush	oru	650		лър	оту	116	655	

Glu	Glu	His	Leu	Leu	Ser	Gly	He	Thr	Asp	Leu	Cys	Leu	Gln	Asp	Leu
			660					665					670		
Pro	Leu	Lys	Glu	Arg	He	Phe	lle	Lys	Leu	Ser	Tyr	Pro	Gln	Asp	Asn
		675					680					685			
Leu	Gln	Pro	Asp	Val	Asn	Leu	Lys	Thr	Leu	Ser	lle	Leu	Ser	Val	Lys
	690					695					700				
Glu	Ser	Cys	lle	Ala	Asn	Ser	Gly	Ser	Asp	Cys	Thr	Ser	His	Leu	Ser
705					710					715					720
Lys	Asp	Leu	Pro	Gly	Ile	Pro	Leu	Gln	Asn	${\tt Glu}$	Ser	Arg	Asp	Ser	Lys
				725					730					735	
Ile	Leu	Lys	Gly	Asp	Gln	Leu	Leu	Gln	Glu	Asp	Tyr	Lys	Val	Asn	Thr
			740					745					750		
Ser	Val	Pro	Tyr	Ser	Val	Ser	Asn	Thr	Val	Val	Lys	Thr	Cys	Asn	Val
		755					760					765			
Arg	Pro	Pro	Asn	Thr	Ala	Leu	Asp	His	Ser	Arg	Lys	Val	Asp	Met	Gln
	770					775					780				
Thr	Thr	Arg	Lys	Ile	Leu	Met	Lys	Lys	Ser	Val	Cys	Leu	Asp	Arg	His
785					790					795					800
Ser	Ser	Asp	Glu	Gln	Ser	Ala	Pro	Val	Phe	Gly	Lys	Ala	Lys	Tyr	Thr
				805					810					815	
Thr	Gln	Arg	Met	Lys	His	Ser	Ser	Gln	Lys	His	Asn	Ser	Ser	His	Phe
			820					825					830		
Lys	Glu	Ser	G] y	His	Asn	Lys	Leu	Ser	Ser	Pro	Lys	He	His	Ile	Lys
		835					840					845			
Glu	Thr	Glu	Gln	Cys	Val	Arg	Ser	Tyr	Glu	Thr	Ala	Glu	Asn	Glu	Glu
	850					855					860				
Ser	Cys	Phe	Pro	Asp	Ser	Thr	Lys	Ser	Ser	Leu	Ser	Ser	Leu	Gln	Cys
865					870					875					880
His	Lys	Lys	Glu	Asn	Asn	Ser	Gly	Thr	Cys	Leu	Asp	Ser	Pro	Leu	Pro
				885					890					895	
Leu	Cys	Gln	Arg	Leu	Lys	Leu	Arg	Phe	Gln	Ser	Thr				
			900					905							

<210> 4738 <211> 637

```
<212> PRT
<213> Homo sapiens
<400> 4738
```

<400)> 47	738													
Met	Leu	Pro	Asn	Phe	Lys	Leu	Tyr	Asn	Phe	He	Glu	He	Phe	Phe	Lys
1				5					10					15	
Pro	Leu	Thr	Pro	Ser	Lys	Asn	Arg	Phe	His	Phe	Val	Ser	Tyr	Phe	Glu
			20					25					30		
Asn	Val	Asn	Phe	Met	Leu	Cys	Trp	Leu	Gln	Glu	Asn	Asn	Phe	Cys	Leu
		35					40					45			•
Leu	Leu	Cys	Phe	Leu	Ser	Gly	Leu	Leu	Ser	Arg	His	Lys	Thr	Lys	Lys
	50					55					60				
Leu	Ser	Ser	Glu	Lys	Asp	He	llis	Glu	lle	Ser	Leu	Ser	Lys	Glu	Ser
65					70					75					80
He	He	Glu	Lys	Ser	Lys	Thr	Leu	Arg	Leu	Lys	G1 y	Ser	lle	Phe	Arg
				85					90					95	
Asn	Glu	Trp	Gln	Asn	Lys	Ser	Glu	Phe	Glu	Gly	Gln	Gln	Gly	Leu	Lys
			100					105					110		
Glu	Arg	Ser	He	Ser	Gln	Lys	Lys	Ile	Val	Ser	Lys	Lys	Met	Ser	Thr
		115					120					125			
Λsp	Arg	Lys	Arg	Pro	Ser	Phe	Thr	Leu	Asn	Gln	Arg	He	His	Asn	Ser
	130					135					140				
Glu	Lys	Ser	Cys	Asp	Ser	His	Leu	Val	Gln	His	G]y	Lys	He	Asp	Ser
145					150					155					160
Asp	Val	Lys	His	Asp	Cys	Lys	Glu	Cys	Gly	Ser	Thr	Phe	Asn	Asn	Val
				165					170					175	
Tyr	Gln	Leu	Thr	Leu	His	Gln	Lys	He	His	Thr	Gly	Glu	Lys	Ser	Cys
			180					185					190		
Lys	Cys	Glu	Lvs	Cys	Gly	Lys		Phe	Ser	His	Ser		Gln	Leu	Thr
		195					200					205			
Leu		G1n	Arg	Phe	His		Gly	Glu	Lys	Pro		Glu	Cys	Gln	Glu
	210					215					220				
	Gly	Lys	Thr	Phe	Thr	Leu	Tyr	Pro	Gln		Asn	Arg	His	Gln	
225					230					235		_		_	240
He	His	Thr	G1 y		Lys	Pro	Tyr	Met			Lys	Cys	Asp		Gly
				245					250					255	

Phe	Phe	Ser	Arg	Leu	Glu	Leu	Thr	Gln	His	Lys	Arg	He	His	Thr	Gly
			260					265					270		
Lys	Lys	Ser	Tyr	Glu	Cys	Lys	Glu	Cys	Gly	Lys	Val	Phe	Gln	Leu	He
		275					280					285			
Phe	Tyr	Phe	Lys	Glu	His	Glu	Arg	lle	His	Thr	Gly	Lys	Lys	Pro	Tyr
	290					295					300				
Glu	Cys	Lys	Glu	Cys	G] y	Lys	Ala	Phe	Ser	Val	Cys	Gly	G1n	Leu	Thr
305					310					315					320
Arg	His	Gln	Lys	He	His	Thr	Gly	Val	Lys	Pro	Tyr	Glu	Cys	Lys	G1u
				325					330					335	
Cys	Gly	Lys	Thr	Phe	Arg	Leu	Ser	Phe	Tyr	Leu	Thr	Glu	His	Arg	Arg
			340					345					350		
Thr	His		Gly	Lys	Lys	Pro		G] u	Cys	Lys	Glu		G1 y	Lys	Ser
		355					360					365			
Phe		Val	Arg	G1 y	Gln		Asn	Arg	His	Lys		He	His	Thr	Gly
	370					375					380		_	_	
	Lys	Pro	Phe	Ala		Lys	Val	Cys	Glu		Ala	Phe	Ser	Tyr	
385					390					395			_	_	400
Gly	Asp	Leu	Arg		His	Ser	Arg	He		Thr	Gly	Glu	Lys		Tyr
0.1			0.1	405	0.1	,		131	410				1	415	6 01
Glu	Cys	Lys		Cys	G1 y	Lys	Ala		Met	Leu	Arg	Ser	Val	Leu	Ihr
C1	11.	C1	420	1	11.	TI.	C1 .	425	1	D	т	C1	430	1	C1.
GIU	HIS	61n	Arg	Leu	HIS	ınr	ыу	val	Lys	Pro	Tyr	GIU	Cys	Lys	GIU
		435					440					445			
Cvc	Cl _v		The	Pho	Ara	Val		Sor	Gln	Па	Sor		His	lve	lve
Cys	450	Lys	1111	1 116	A1 g	455		261	OIII	116	460	Leu	1113	Lys	Lys
ماآ		Thr	Asn	Val	lve			lve	Cve	Val		Cvs	Gly	lve	Thr
465	1115	1111	пор	, (1)	470	110	.,.	15,5	0,5	475	m s	0,5	01,	Ly S	480
	Arg	Phe	GIv	Phe		Len	Thr	61u	His		Arg	He	His	Thr	
1110	111 8	1110	91,	485	* . *	1200			490	5211	5		312.0	495	~
Glu	Lvs	Pro	Tvr		Cvs	Lvs	Glu	Cvs		Lvs	Ala	Phe	lle		Are
0,0	.,, .,		500		.,,			505	,	22, 4.			510	0	2
Glv	Asn	Leu		Glu	His	Leu	Lvs		His	Ser	G1 v	Leu	Lys	Pro	Tvr
3		515					520				3	525	,		
Asn	Cve		Glu	Cve	Glv	lve		Phe	Ser	Aro	Arg		Gln	Phe	Thr

Glu His Gln Lys Ile His Thr Gly Val Lys Pro Tyr Lys Cys Lys Glu Cvs Glv Lys Ala Phe Ser Arg Ser Val Asp Leu Arg Ile His Gln Arg lle His Thr Gly Glu Lys Pro Tyr Glu Cys Lys Gln Cys Gly Lys Ala Phe Arg Leu Asn Ser His Leu Thr Glu His Gln Arg Ile His Thr Gly Glu Lys Pro Tyr Glu Cys Lys Val Cys Arg Lys Ala Phe Arg Gln Tyr Ser His Leu Tyr Gln His Gln Lys Thr His Asn Val Ile

<210> 4739

<211> 170

<212> PRT

<213> Homo sapiens

⟨400⟩ 4739

Met Arg Ala Ala Val Pro His Gln Gly 11e Lys Phe Phe Lys Thr Leu Val Pro Tyr Asp Glu Gly Ala Pro Ala Phe Glu Gly Arg Ala Pro Ala Phe Ser His Ala Ser Leu His Pro Arg Tyr Asp Leu Met Tyr Gln Cys Trp Ser Ala Asp Pro Lys Gln Arg Pro Ser Phe Thr Cys Leu Arg Met

Glu Leu Glu Asn 11e Leu Gly Gln Leu Ser Val Leu Ser Ala Ser Gln

Asp Pro Leu Tyr Ile Asn Ile Glu Arg Ala Glu Glu Pro Thr Ala Gly

Gly Ser Leu Glu Leu Pro Gly Arg Asp Gln Pro Tyr Ser Gly Ala Gly

Asp Gly Ser Gly Met Gly Ala Val Gly Gly Thr Pro Ser Asp Cys Arg

Tyr 11e Leu Thr Pro Gly Gly Leu Ala Glu Gln Pro Gly Gln Ala Glu His Gln Pro Glu Ser Pro Leu Asn Glu Thr Gln Arg Leu Leu Leu Gln Gln Gly Leu Leu Pro His Ser Ser Cys <210> 4740 <211> 492 <212> PRT <213> Homo sapiens <400> 4740 Met Ala Ser Pro Ser Gly Lys Gly Ala Arg Ala Leu Glu Ala Pro Gly Cys Gly Pro Arg Pro Leu Ala Arg Asp Leu Val Asp Ser Val Asp Asp Ala Glu Gly Leu Tyr Val Ala Val Glu Arg Cys Pro Leu Cys Asn Thr Thr Arg Arg Arg Leu Thr Cys Ala Lys Cys Val Gln Ser Gly Asp Phe Val Tyr Phe Asp Gly Arg Asp Arg Glu Arg Phe lle Asp Lys Lys Glu Arg Leu Ser Arg Leu Lys Ser Lys Gln Glu Glu Phe Gln Lys Glu Val Leu Lys Ala Met Glu Gly Lys Trp Ile Thr Asp Gln Leu Arg Trp Lys lle Met Ser Cys Lys Met Arg lle Glu Gln Leu Lys Gln Thr lle Cys Lys Gly Asn Glu Glu Met Glu Lys Asn Ser Glu Gly Leu Leu Lys Thr Lys Glu Lys Asn Gln Lys Leu Tyr Ser Arg Ala Gln Arg His Gln Glu

Lys Lys Glu Lys Ile Gln Arg His Asn Arg Lys Leu Gly Asp Leu Val

				165					170					175	
Glu	Lys	Lys	Thr	He	Asp	Leu	Arg	Ser	His	Tyr	Glu	Arg	Leu	Ala	Asn
			180					185					190		
Leu	Arg	Arg	Ser	His	He	Leu	Glu	Leu	Thr	Ser	Val	lle	Phe	Pro	He
		195					200					205			
Glu	Glu	Val	Lys	Thr	Gly	Val	Arg	Asp	Pro	Ala	Asp	Val	Ser	Ser	Glu
	210			•		215					220				
Ser	Asp	Ser	Ala	Met	Thr	Ser	Ser	Thr	Val	Ser	Lys	Leu	Ala	Glu	Ala
225					230					235					240
Arg	Arg	Thr	Thr	Tyr	Leu	Ser	Gly	Arg	Trp	Val	Cys	Asp	Asp	His	Asn
				245					250					255	
Gly	Asp	Thr	Ser	lle	Ser	Ile	Thr	Gly	Pro	Trp	lle	Ser	Leu	Pro	Asn
			260					265					270		
Asn	Gly	Asp	Tyr	Ser	Ala	Tyr	Tyr	Ser	Trp	Val	Glu	Glu	Lys	Lys	Thr
		275					280					285			
Thr	Gln	Gly	Pro	Asp	Met	Glu	Gln	Ser	Asn	Pro	Ala	Tyr	Thr	He	Ser
	290					295					300				
A 3	A 1 a	Lou	Cvc	Tyr	Ala	Thr	Gln	Leu	Val	Asn	He	Leu	Ser	His	He
Ala	Ala	Leu	Cys	1) 1	MIG	1 111	V	150 0	,				~ ~ .		
305	піа	Leu	Cys	1 9 1	310	1111	· · · ·	,,,,,	, 41	315					320
305					310					315				Cys	320
305					310					315					320
305 Leu	Asp	Val	Asn	Leu 325	310 Pro	Lys	Lys	Leu	Cys 330	315 Asn	Ser	Glu	Phe	Cys	320 Gly
305 Leu	Asp	Val	Asn	Leu 325	310 Pro	Lys	Lys	Leu	Cys 330	315 Asn	Ser	Glu	Phe	Cys 335	320 Gly
305 Leu Glu	Asp Asn	Val Leu	Asn Ser 340	Leu 325 Lys	310 Pro Gln	Lys Lys	Lys Phe	Leu Thr 345	Cys 330 Arg	315 Asn Ala	Ser Val	G1u Lys	Phe Lys 350	Cys 335	320 Gly Asn
305 Leu Glu	Asp Asn	Val Leu	Asn Ser 340	Leu 325 Lys	310 Pro Gln	Lys Lys	Lys Phe	Leu Thr 345	Cys 330 Arg	315 Asn Ala	Ser Val	G1u Lys	Phe Lys 350	Cys 335 Leu	320 Gly Asn
305 Leu Glu Ala	Asp Asn Asn	Val Leu Ile 355	Asn Ser 340 Leu	Leu 325 Lys Tyr	310 Pro Gln Leu	Lys Lys Cys	Lys Phe Phe 360	Leu Thr 345 Ser	Cys 330 Arg Gln	315 Asn Ala His	Ser Val	Glu Lys Asn 365	Phe Lys 350 Leu	Cys 335 Leu	320 Gly Asn Gln
305 Leu Glu Ala	Asp Asn Asn	Val Leu Ile 355	Asn Ser 340 Leu	Leu 325 Lys Tyr	310 Pro Gln Leu	Lys Lys Cys	Lys Phe Phe 360 Arg	Leu Thr 345 Ser	Cys 330 Arg Gln	315 Asn Ala His	Ser Val	Glu Lys Asn 365 Leu	Phe Lys 350 Leu	Cys 335 Leu Asp	320 Gly Asn Gln
305 Leu Glu Ala Leu	Asp Asn Asn Gln 370	Val Leu lle 355 Pro	Asn Ser 340 Leu Leu	Leu 325 Lys Tyr His	310 Pro Gln Leu Thr	Lys Lys Cys Leu 375	Lys Phe Phe 360 Arg	Leu Thr 345 Ser Asn	Cys 330 Arg Gln Leu	315 Asn Ala His	Ser Val Val Tyr 380	Glu Lys Asn 365 Leu	Phe Lys 350 Leu Val	Cys 335 Leu Asp	320 Gly Asn Gln Pro
305 Leu Glu Ala Leu	Asp Asn Asn Gln 370	Val Leu lle 355 Pro	Asn Ser 340 Leu Leu	Leu 325 Lys Tyr His	310 Pro Gln Leu Thr	Lys Lys Cys Leu 375	Lys Phe Phe 360 Arg	Leu Thr 345 Ser Asn	Cys 330 Arg Gln Leu	315 Asn Ala His	Ser Val Val Tyr 380	Glu Lys Asn 365 Leu	Phe Lys 350 Leu Val	Cys 335 Leu Asp	320 Gly Asn Gln Pro
305 Leu Glu Ala Leu Ser 385	Asn Asn Gln 370 Ser	Val Leu Ile 355 Pro Glu	Asn Ser 340 Leu Leu	Leu 325 Lys Tyr His	310 Pro Gln Leu Thr Gly 390	Lys Lys Cys Leu 375 Arg	Lys Phe Phe 360 Arg	Leu Thr 345 Ser Asn	Cys 330 Arg Gln Leu Pro	315 Asn Ala His Met Phe 395	Ser Val Val Tyr 380 Glu	Glu Lys Asn 365 Leu Val	Phe Lys 350 Leu Val	Cys 335 Leu Asp	320 Gly Asn Gln Pro Asp 400
305 Leu Glu Ala Leu Ser 385	Asn Asn Gln 370 Ser	Val Leu Ile 355 Pro Glu	Asn Ser 340 Leu Leu	Leu 325 Lys Tyr His	310 Pro Gln Leu Thr Gly 390	Lys Lys Cys Leu 375 Arg	Lys Phe Phe 360 Arg	Leu Thr 345 Ser Asn	Cys 330 Arg Gln Leu Pro	315 Asn Ala His Met Phe 395	Ser Val Val Tyr 380 Glu	Glu Lys Asn 365 Leu Val	Phe Lys 350 Leu Val	Cys 335 Leu Asp Ser	320 Gly Asn Gln Pro Asp 400
305 Leu Glu Ala Leu Ser 385 Leu	Asn Asn Gln 370 Ser	Val Leu Ile 355 Pro Glu Glu	Asn Ser 340 Leu Leu His	Leu 325 Lys Tyr His Leu Met 405	310 Pro Gln Leu Thr Gly 390 Glu	Lys Lys Cys Leu 375 Arg	Lys Phe Phe 360 Arg Ser	Leu Thr 345 Ser Asn Gly	Cys 330 Arg Gln Leu Pro Pro 410	315 Asn Ala His Met Phe 395 Gly	Ser Val Val Tyr 380 Glu Val	Glu Lys Asn 365 Leu Val	Phe Lys 350 Leu Val Arg	Cys 335 Leu Asp Ser Ala	320 Gly Asn Gln Pro Asp 400 Ser
305 Leu Glu Ala Leu Ser 385 Leu	Asn Asn Gln 370 Ser	Val Leu Ile 355 Pro Glu Glu	Asn Ser 340 Leu Leu His	Leu 325 Lys Tyr His Leu Met 405	310 Pro Gln Leu Thr Gly 390 Glu	Lys Lys Cys Leu 375 Arg	Lys Phe Phe 360 Arg Ser	Leu Thr 345 Ser Asn Gly	Cys 330 Arg Gln Leu Pro Pro 410	315 Asn Ala His Met Phe 395 Gly	Ser Val Val Tyr 380 Glu Val	Glu Lys Asn 365 Leu Val	Phe Lys 350 Leu Val Arg	Cys 335 Leu Asp Ser Ala Glu 415	320 Gly Asn Gln Pro Asp 400 Ser
305 Leu Glu Ala Leu Ser 385 Leu Asp	Asp Asn Asn Gln 370 Ser Glu	Val Leu Ile 355 Pro Glu Glu Ser	Asn Ser 340 Leu His Ser Gly 420	Leu 325 Lys Tyr His Leu Met 405 Asp	310 Pro Gln Leu Thr Gly 390 Glu Glu	Lys Cys Leu 375 Arg Phe	Lys Phe Phe 360 Arg Ser Val	Leu Thr 345 Ser Asn Gly Asp Ser 425	Cys 330 Arg Gln Leu Pro Pro 410 Asp	315 Asn Ala His Met Phe 395 Gly Glu	Ser Val Val Tyr 380 Glu Val	Glu Lys Asn 365 Leu Val Ala	Phe Lys 350 Leu Val Arg Gly Asp 430	Cys 335 Leu Asp Ser Ala Glu 415	320 Gly Asn Gln Pro Asp 400 Ser Gly
305 Leu Glu Ala Leu Ser 385 Leu Asp	Asp Asn Asn Gln 370 Ser Glu	Val Leu Ile 355 Pro Glu Glu Ser	Asn Ser 340 Leu His Ser Gly 420	Leu 325 Lys Tyr His Leu Met 405 Asp	310 Pro Gln Leu Thr Gly 390 Glu Glu	Lys Cys Leu 375 Arg Phe	Lys Phe Phe 360 Arg Ser Val	Leu Thr 345 Ser Asn Gly Asp Ser 425	Cys 330 Arg Gln Leu Pro Pro 410 Asp	315 Asn Ala His Met Phe 395 Gly Glu	Ser Val Val Tyr 380 Glu Val	Glu Lys Asn 365 Leu Val Ala	Phe Lys 350 Leu Val Arg Gly Asp 430	Cys 335 Leu Asp Ser Ala Glu 415 Leu	320 Gly Asn Gln Pro Asp 400 Ser Gly

Ile Ala Ser Ser Ser Ala Gly Gly Met Ile Ser Ser Ala Ala Ala Ser Val Thr Ser Trp Phe Lys Ala Tyr Thr Gly His Arg <210> 4741 <211> 415 <212> PRT <213> Homo sapiens <400> 4741 Met Leu Ser Met Thr Tyr Ser Glu Ser Leu Arg Ser Val Ser Ser Arg

Cys His Ser Glu Trp Ala Leu His Pro Val Arg Gln Thr Asp Thr Leu Glu Leu Gln Arg Leu Arg Glu Val Arg Ala Ala Ala Gln Ala Arg Asn Met Glu Ser Phe Leu Arg Met His Gly Leu Ser Leu Asp Gly Cys Thr Ala Gln Arg Thr Gly Met Lys Tyr Arg Asn Leu Gly Lys Ser Gly Leu Arg Val Ser Cys Leu Gly Leu Gly Thr Trp Val Thr Phe Gly Gly Gln Ile Thr Asp Glu Met Ala Glu Gln Leu Met Thr Leu Ala Tyr Asp Asn Gly lle Asn Leu Phe Asp Thr Ala Glu Val Tyr Ala Ala Gly Lys Ala Glu Val Val Leu Gly Asn Ile Ile Lys Lys Lys Gly Trp Arg Arg Ser Ser Leu Val Ile Thr Thr Lys Ile Phe Trp Gly Gly Lys Ala Glu Thr Glu Arg Gly Leu Ser Arg Lys His 11e 11e Glu Gly Leu Lys Ala Ser

Leu Glu Arg Leu Gln Leu Glu Tyr Val Asp Val Val Phe Ala Asn Arg

			180					185					190		
Pro	Asp	Pro	Asn	Thr	Pro	Met	Glu	G1 y	Asp	Pro	Phe	Ser	Ser	Ser	Lys
		195					200					205			
Ser	Arg	Thr	Phe	He	lle	Glu	Glu	Thr	Val	Arg	Ala	Met	Thr	His	Val
	210					215					220				
He	Asn	Gln	Gly	Met	Ala	Met	Tyr	Trp	Gly	Thr	Ser	Arg	Trp	Ser	Ser
225					230					235					240
Met	Glu	lle	Met	Glu	Ala	Tyr	Ser	Val	Ala	Arg	Gln	Phe	Asn	Leu	Thr
				245					250					255	
Pro	Pro	lle	Cys	Glu	G1n	Ala	Glu	Tyr	His	Met	Phe	Gln	Arg	Glu	Lys
			260					265					270		
Val	Glu	Val	Gln	Leu	Pro	Glu	Leu	Phe	His	Lys	lle	Gly	Val	Gly	Ala
		275					280					285			
Met	Thr	Trp	Ser	Pro	Leu	Ala	Cys	Gly	He	Va]	Ser	Gly	Lys	Tyr	Asp
	290					295					300				
Ser	Gly	lle	Pro	Pro	Tyr	Ser	Arg	Ala	Ser	Leu	Lys	Gly	Tyr	Gln	Trp
305					310					315					320
Leu	Lys	Asp	Lys	Ile	Leu	Ser	Glu	Glu	Gly	Arg	Arg	Gln	Gln	Ala	Lys
				325					330					335	
Leu	Lys	Glu	Leu	Gln	Ala	He	Ala	Glu	Arg	Leu	Gly	Cys	Thr	Leu	Pro
			340					345					350		
Gln	Leu	Ala	lle	Ala	Trp	Cys	Leu	Arg	Asn	Glu	Gly	Val	Ser	Ser	Val
		355					360					365			
Leu	Leu	Gly	Ala	Ser	Asn	Ala	Asp	Gln	Leu	Met	Glu	Asn	lle	Gly	Ala
	370					375					380				
He	Gln	Val	Leu	Pro	Lys	Leu	Ser	Ser	Ser	lle	lle	His	Glu	lle	Asp
385					390					395					400
Ser	11e	Leu	Gly	Asn	Lys	Pro	Tyr	Ser	Lys	Lys	Asp	Tyr	Arg	Ser	
				405					410					415	

<210> 4742

<211> 127

<212> PRT

<213> Homo sapiens

<400> 4742 Met Arg Gln Trp Pro Val Ala Val Pro Cys Ser Arg Gln Gln Cys Cys 10 15 Arg Ser Ser Leu Gln Pro Thr Ile Cys His Tyr Phe Met Arg Leu Leu 20 25 Lys Asp Lys Gly Leu Leu Leu Arg Cys Tyr Thr Gln Asn Ile Asp Thr 40 45 Leu Glu Arg Ile Ala Gly Leu Glu Gln Glu Asp Leu Val Glu Ala His 50 55 60 Gly Thr Phe Tyr Thr Ser His Cys Val Ser Ala Ser Cys Arg His Glu 70 75 Tyr Pro Leu Ser Trp Met Lys Glu Lys Ile Phe Ser Glu Val Thr Pro 90 Lys Cys Glu Asp Cys Gln Ser Leu Val Lys Pro Gly Glu Pro Leu Ala 105 Arg Asp Leu Pro Arg Trp Ile Trp Val Gly Val Pro Ser Ser Pro 120 125

<210> 4743

<211> 540

<212> PRT

<213> Homo sapiens

<400> 4743

Met Pro Asn Lys Asn Lys Lys Glu Lys Glu Ser Pro Lys Ala Gly Lys

1 5 10 15

Ser Gly Lys Ser Ser' Lys Glu Gly Gln Asp Thr Val Glu Ser Glu Gln

Ser Gly Lys Ser Ser Lys Glu Gly Gln Asp Thr Val Glu Ser Glu Gln
20 25 30

lle Ser Val Arg Lys Asn Ser Leu Val Ala Val Pro Ser Thr Val Ser

Ala Lys Ile Lys Val Pro Val Ser Gln Pro Ile Val Lys Lys Asp Lys
50 55 60

Arg Gln Asn Ser Ser Arg Phe Ser Ala Ser Asn Asn Arg Glu Leu Gln
65 70 75 80

Lys	Leu	Pro	Ser	Leu	Lys	Asp	Val	Pro	Pro	Ala	Asp	Gln	Glu	Lys	Leu
				85					90					95	
Phe	He	Gln	Lys	Leu	Arg	Gln	Cys	Cys	Val	Leu	Phe	Asp	Phe	Val	Ser
			100					105					110		
Asp	Pro	Leu	Ser	Asp	Leu	Lys	Trp	Lys	Glu	Val	Lys	۸rg	Ala	Ala	Leu
		115					120					125			
Ser	Glu	Met	Val	Glu	Tyr	Пе	Thr	His	Asn	Arg	Asn	Val	lle	Thr	Glu
	130					135					140				
Pro	lle	Tyr	Pro	Glu	Val	Val	His	Met	Phe	Ala	Val	Asn	Met	Phe	Arg
145					150					155					160
Thr	Leu	Pro	Pro	Ser	Ser	Asn	Pro	Thr	Gly	Ala	Glu	Phe	Asp	Pro	Glu
				165					170					175	
Glu	Asp	Glu	Pro	Thr	Leu	Glu	Ala	Ala	Trp	Pro	His	Leu	Gln	Leu	Val
			180					185					190		
Tyr	Glu	Phe	Phe	Leu	Arg	Phe	Leu	Glu	Ser	Pro	Asp	Phe	Gln	Pro	Asn
		195					200					205			
He	Ala	Lys	Lys	Tyr	lle	Asp	Gln	Lys	Phe	Val	Leu	Gln	Leu	Leu	Glu
	210					215					220				
Leu	Phe	Asp	Ser	Glu	Asp	Pro	Arg	Glu	Arg	Asp	Phe	Leu	Lys	Thr	Thr
225					230					235					240
Leu	His	Arg	Ile	Tyr	Gly	Lys	Phe	Leu	Gly	Leu	Arg	Ala	Tyr	Ile	Arg
				245					250					255	
Lys	Gln	lle	Asn	Asn	lle	Phe	Tyr	Arg	Phe	He	Tyr	Glu	Thr	Glu	His
			260					265					270		
His	Asn	Gly	He	Ala	Glu	Leu	Leu	Glu	lle	Leu	Gly	Ser	He	He	Asn
		275					280					285			
Gly	Phe	Ala	Leu	Pro	Leu	Lys	Glu	Glu	His	Lys	lle	Phe	Leu	Leu	Lys
	290					295					300				
Val	Leu	Leu	Pro	Leu	His	Lys	Va1	Lys	Ser	Leu	Ser	Val	Tyr	His	Pro
305					310					315					320
Gln	Leu	Ala	Tyr	Cys	Val	Val	Gln	Phe	Leu	Glu	Lys	Asp	Ser	Thr	Leu
				325					330					335	
Thr	G]u	Pro	Val	Val	Met	Ala	Leu	Leu	Lys	Tyr	Trp	Pro	Lys	Thr	His
			340					345					350		
Ser	Pro	Lys	Glu	Val	Met	Phe	Leu	Asn	Glu	Leu	Glu	Glu	He	Leu	Asp
		355					360					365			

Val Ile Glu Pro Ser Glu Phe Val Lys Ile Met Glu Pro Leu Phe Arg Gln Leu Ala Lys Cys Val Ser Ser Pro His Phe Gln Val Ala Glu Arg Ala Leu Tyr Tyr Trp Asn Asn Glu Tyr Ile Met Ser Leu Ile Ser Asp Asn Ala Ala Lys Ile Leu Pro lle Met Phe Pro Ser Leu Tyr Arg Asn Ser Lys Thr His Trp Asn Lys Thr Ile His Gly Leu Ile Tyr Asn Ala Leu Lys Leu Phe Met Glu Met Asn Gln Lys Leu Phe Asp Asp Cys Thr Gln Gln Phe Lys Ala Glu Lys Leu Lys Glu Lys Leu Lys Met Lys Glu Arg Glu Glu Ala Trp Val Lys Ile Glu Asn Leu Ala Lys Ala Asn Pro Gln Ala Gln Lys Asp Pro Lys Lys Asp Arg Pro Leu Ala Arg Arg Lys Ser Glu Leu Pro Gln Asp Pro His Thr Lys Lys Ala Leu Glu Ala His Cys Arg Ala Asp Glu Leu Ala Ser Gln Asp Gly Arg

<210> 4744

<211> 361

<212> PRT

<213> Homo sapiens

<400> 4744

Leu		Lys	Glu	Glu	Phe		His	Val	lle	Ala		Gln	Gln	Gly	Glu
	50					55					60				
	Gly	Glu	Phe	Val		Lys	Asp	Pro	Asp		Val	Ala	Arg	Leu	
65					70					75					80
Gly	Arg	Arg	Lys	Cys	Lys	Pro	Gln	Met	Asn	Tyr	Asp	Lys	Leu		Arg
				85					90					95	
Ala	Leu	Arg	Tyr	Tyr	Tyr	Asn	Lys	Arg	Ile	Leu	His	Lys	Thr	Lys	Gly
			100					105					110		
Lys	Arg	Phe	Thr	Tyr	Lys	Phe	Asn	Phe	Ser	Lys	Leu	Пe	Val	Val	Asn
		115					120					125			
Tyr	Pro	Leu	Trp	Glu	Val	Arg	Ala	Pro	Pro	Ser	Pro	His	Leu	Leu	Leu
	130					135					140				
Gly	Ala	Pro	Ala	Leu	Cys	Arg	Pro	Ala	Leu	Val	Pro	Val	Gly	Val	Gln
145					150					155					160
Ser	Glu	Leu	Leu	His	Ser	Met	Leu	Phe	Ala	His	Gln	Ala	Met	Val	Glu
				165					170					175	
Gln	Leu	Thr	G1 y	Gln	Gln	Thr	Pro	Arg	Gly	Pro	Pro	Glu	Thr	Ser	Gly
			180					185					190		
Asp	Lys	Lys	Gly	Ser	Ser	Ser	Ser	Val	Tyr	Arg	Leu	Gly	Ser	Ala	Pro
		195					200					205			
Gly	Pro	Cys	Arg	Leu	Gly	Leu	Cys	Cys	His	Leu	Gly	Ser	Val	Gln	Gly
	210					215					220				
Glu	Leu	Pro	Gly	Val	Ala	Ser	Phe	Thr	Pro	Pro	Leu	Pro	Pro	Pro	Leu
225					230					235					240
Pro	Ser	Asn	Trp	Thr	Cys	Leu	Ser	Gly	Pro	Phe	Leu	Pro	Pro	Leu	Pro
				245					250					255	
Ser	Glu	Gln	Gln	Leu	Pro	Gly	Ala	Phe	Lys	Pro	Asp	He	Leu	Leu	Pro
			260					265					270		
Gly	Pro	Arg	Ser	Leu	Pro	Gly	Ala	Trp	His	Phe	Pro	Gly	Leu	Pro	Leu
		275					280					285			
Leu	Ala	Gly	Leu	Gly	Gln	Gly	Ala	Gly	Glu	Arg	Leu	Trp	Leu	Leu	Ser
	290					295					300				
Leu	Arg	Pro	Glu	Gly	Leu	Glu	Val	Lys	Pro	Ala	Pro	Met	Met	Glu	Ala
305	J			•	310			-		315					320
	Glv	Glv	Leu	Asp		Arg	Glu	Val	Phe		Pro	Glu	Thr	Arg	
,	- ,	J		325		3			330	•				335	J

Leu Lys Thr Gly Glu Glu Ser Leu Thr Ser Pro Asn Leu Glu Asn Leu 340 345 350

Lys Ala Val Trp Pro Leu Asp Pro Pro 355 360

<210> 4745

<211> 169

<212> PRT

<213> Homo sapiens

<400> 4745

Met Met Lys Thr Leu Asn Ile Leu Asp Met Glu Gly Met Phe Leu Asn
1 5 10 15

Thr Ile Lys Ala lle Tyr Asp Lys Leu lle Ala Asn Ile Val Leu Ser 20 25 30

Gly Lys Lys Leu Lys Ala Phe Pro Ile Arg Pro Arg Thr Arg Gln Gly
35 40 45

Cys Pro Leu Leu Pro Leu Leu Phe Asn Ile Val Pro Glu Val Leu Ala 50 55 60

Arg Ala Ile Arg Gln Glu Lys Glu Ile Lys Asp Ile Gln Ile Gly Lys
65 70 75 80

Ser Glu Met Lys Leu Ser Leu Phe Ala Asp Asn Met 11e Leu His 11e 85 90 95

Glu Asn Leu Lys Asp Cys Thr Lys Thr Pro Phe Glu Leu Asp Lys Ser 100 105 110

Ser Lys Val Ala Gly Tyr Lys 11e Asn Leu Gln Lys Ser Val His Phe 115 · 120 125

Tyr Thr Leu Thr Val Asn Phe Leu Lys Lys Lys Leu Arg Lys Gln Ser 130 135 140

His Leu Gln Tyr His Leu Leu Leu Ser Lys IIe Leu Arg Arg Lys Phe 145 150 155 160

Asn Gln Gly Ser Glu Arg Ser Ile His

```
<210> 4746
<211> 182
<212> PRT
<213> Homo sapiens
<400> 4746
Met Trp Ile Leu Arg Val Leu Leu Gly Arg His Leu Pro Gln Trp Asn
                                     10
Ala Ser Ser Glu Trp Cys Met Ala Gly Pro His Gly Gly Ser Thr Gln
             20
                                 25
                                                      30
Gly Leu Asn Thr Gly Lys Glu Leu Ala Arg Gly Val Arg Thr Ser Gly
                             40
lle Trp Glu Ser Ala Ala Trp Asp His Lys Glu Ser Lys Ser Ser Ala
                         55
Thr Cys Trp Ser Leu Arg Pro Glu Gly Arg Pro Asp Phe Arg Asp His
 65
                     70
                                          75
Pro Trp Met Ser Leu Asp Trp Glu Leu Tyr Val Asp Gly Ser Asn Phe
                 85
                                     90
Val Asn Ser Gln Gly Glu Arg Cys Val Gly Tyr Ala Val Val Thr Leu
                                                     110
            100
                                 105
Asp Ala Val lle Glu Ala Lys Ser Leu Pro Gln Gly Thr Ser Ala Gln
                            120
                                                 125
Lys Ala Glu Leu lle Ala Leu lle Arg Ala Ser Glu Leu Ser Glu Gly
                        135
Lys Thr Val Asn lle Tyr Thr Asp Ser Gln Tyr Ala Phe Leu Thr Leu
                                                              160
145
                    150
                                         155
Gln Val His Gly Ala Leu Tyr Lys Glu Lys Gly Leu Leu Asn Ser Gly
                165
                                     170
                                                         175
Gly Lys Asp Val Lys Tyr
```

<210> 4747

<211> 957

<212> PRT

<213> Homo sapiens

<400	> 47	747													
Met	Thr	Ser	Phe	Leu	Lys	Pro	Glu	Asn	Ala	Leu	Lys	Arg	Ala	Glu	Glu
1				5					10					15	
Leu	He	Asn	Val	Gly	Gln	Lys	Gln	Asp	Ala	Leu	Gln	Thr	Leu	His	Asp
			20					25					30		
Leu	He	Thr	Ser	Lys	Arg	Tyr	Arg	Ala	Trp	Gln	Lys	Thr	Leu	Glu	Arg
		35					40					45			
He	Met	Phe	Lys	Tyr	Val	Glu	Leu	Cys	Val	Asp	Met	Arg	Lys	Gly	Arg
	50					55					60				
Phe	Ala	Lys	Asp	G1 y	Leu	He	Gln	Tyr	Arg	Ile	lle	Cys	Gln	Gln	Val
65					70					75					80
Asn	Val	Ser	Ser	Leu	Glu	Glu	Val	He	Lys	His	Phe	Met	Gln	Leu	Ser
				85					90					95	
Thr	Glu	Lys	Ala	Glu	Gln	Ala	Arg	Ser	Gln	Λla	Gln	Ala	Leu	Glu	Glu
			100					105					110		
Ala	Leu	Asp	Val	Asp	Asp	Leu	Glu	Ala	Asp	Lys	Arg	Pro	Glu	Asp	Leu
		115					120					125			
Met		Ser	Tyr	Val	Ser		Glu	Lys	Gly	Lys		Arg	Ser	Asp	Arg
	130					135					140				
Glu	Thr	Val	Thr	Pro		Phe	Lys	Phe	Leu		Glu	Thr	Tyr	Arg	
145					150					155					160
Val	Leu	Glu	lle		Arg	Asn	Asn	Ser		Leu	GIu	Ala	Leu		Ala
				165		151	0.1	151	170		0.1	Tr.		175	T)
Met	Thr	Ala		Arg	Ala	Phe	Gin		Cys	Lys	Gln	lyr	Lys	Arg	Ihr
m	0.1		180			0	61	185	11.	Δ.	Δ	117 -	190	A 1 -	Λ
Ihr	Glu			Arg	Leu	Cys				Arg	Asn	111 S 205	Leu	АТА	ASN
,	4	195		A	Λ	C1		A		Dwa	A on	200		410	Dro
Leu		Lys	ıyr	Arg	Asp		Arg	Asp	Arg	rro		Leu	Ser	Ala	710
C1	210	1	C1	ارما	Т	215	Aan	Thu	Ana	Dho	220	Cln	Lou	Lvc	По
	261	Leu	GIII	Leu	230	Leu	nsp	1111	MIG	235	Glu	0111	Leu	Lys	240
225	The	Clu	Lou	C1v		Tro	Cln	Clu	Ala		Δra	Sor	Val	Glu	
wia	1 1111.	oru	เซน	245	ren	11 b	0111	01u	250		Λ1 g	261	val	255	nsp
110	Hic	Clv	ا من		Cvc	ىرم [Val	lve			Pro	lve	Pro		l en
, 16	1113	UIY	260		Oya	Leu		265		4113	.10	درر	270		204
			200					200							

Met	Val	Val	Tyr	Tyr	Val	Lys	Leu	Thr	Glu	Пe	Phe	Trp	Ile	Ser	Ser
		275					280					285			
Ser	llis	Leu	Tyr	His	Ala	Tyr	Ala	Trp	Phe	Lys	Leu	Phe	Leu	Leu	Gln
	290					295					300				
Lys	Ser	Phe	Asn	Lys	Asn	Leu	Ser	Gln	Lys	Asp	Leu	Gln	Leu	Ile	Ala
305					310					315					320
Ser	Ser	Val	Val	Leu	Ala	Ala	Leu	Ser	Val	Pro	Pro	llis	Asp	Arg	Thr
				325					330					335	
His	Gly	Ala	Ser	His	Leu	Glu	Leu	Glu	His	Glu	Lys	Glu	Arg	Asn	Leu
			340					345					350		
Arg	Met	Ala	Asn	Leu	Ile	Gly	Phe	Asn	Leu	Glu	Thr	Lys	Pro	Glu	Ser
		355					360					365			
Arg	Glu	Met	Leu	Ser	Arg	Ala	Ser	Leu	Leu	Λla	Glu	Leu	Ala	Ser	Lys
	370					375					380				
Gly	Val	Met	Ser	Cys	Val	Thr	Gln	Glu	Val	Lys	Asp	He	Tyr	His	Leu
385					390					395					400
Leu	Glu	His	Glu	Phe	Tyr	Pro	Ser	Asp	Leu	Ala	Leu	Lys	Ala	Leu	Pro
				405					410					415	
Leu	lle	Thr	Lys	Πle	Ser	Lys	Leu	Gly	Gly	Lys	Leu	Ser	Thr	Ala	Ser
			420					425					430		
Ser	Val	Pro	Glu	Val	Gln	Leu	Ala	Gln	Tyr	Val	Pro	Ala	Leu	Glu	Arg
		435					440					445			
Leu	Ala	Thr	Met	Arg	Leu	Leu	Gln	Gln	Val	Ser	Asn	Val	Tyr	Gln	Ser
	450					455					460				
Met	Lys	He	G]u	Thr	Leu	Ser	Gly	Met	He	Pro	Phe	Phe	Asp	Phe	Ala
465					470					475					480
Gln	Va]	G] u	Lys	lle	Ser	Val	Asp	Ala	Val	Lys	Gln	Lys	Phe	Val	Ser
				485					490					495	
Met	Lys	Val	Asp	His	Met	Lys	Asn	Ala	Val	11e	Phe	Ser	Lys	Lys	Ser
			500					505					510		
Leu	Glu	Ser	Asp	Gly	Leu	Arg	Asp	His	Leu	G1 y	Asn	Phe	Ala	Glu	Gln
		515					520					525			
Leu	Asn	Lys	Ala	Arg	Gln	Met	He	Tyr	Pro	Pro	Asp	Gly	Arg	Pro	Ser
	530					535					540				
Lys	Leu	G1y	Ala	Leu	Leu	Pro	Thr	Leu	Thr	Glu	Val	Val	Ala	Lys	Glu
545					550					555					560

His	Lys	Arg	Leu		Ala	Arg	Lys	Ser		He	Glu	Lys	Arg		Glu
				565					570					575	
Glu	Gln	Glu	Arg	GIn	Leu	Leu	Glu		Glu	Arg	Glu	Glu		Ser	Lys
			580					585					590		
Arg	Leu	Arg	Leu	GIn	Lys	Пe	Thr	Glu	Glu	Ala	Glu	Gln	Arg	Arg	Leu
		595					600					605			
Ala	Thr	Glu	Tyr	Glu	Gln	Arg	Lys	Asn	Gln	Arg	He	Leu	Arg	Glu	Ile
	610					615					620				
Glu	G]u	Arg	Glu	Asn	Glu	Glu	Ala	Gln	Ala	Leu	Leu	Gln	Glu	Ala	Glu
625					630					635					640
Lys	Arg	He	Lys	Lys	Lys	Gly	Lys	Lys	Pro	lle	He	Glu	Gly	Asp	Lys
				645					650					655	
He	Thr	Lys	Gln	Thr	Leu	Met	Glu	Leu	Thr	Leu	Thr	Glu	Gln	Leu	Arg
			660					665					670		
Glu	Arg	Gln	Glu	Met	Glu	Lys	Lys	Leu	Gln	Lys	Leu	Ala	Lys	Thr	Met
		675					680					685			
Asp	Tyr	Leu	Glu	Arg	Ala	Lys	Arg	Glu	Glu	Ala	Ala	Pro	Leu	lle	Glu
	690					695					700				
Ala	Ala	Tyr	Gln	Gln	Arg	Leu	Val	Glu	Glu	Arg	Leu	Leu	His	Glu	Arg
705					710					715					720
Glu	Gln	Gln	G1n	Glu	Val	Glu	Leu	Ser	Lys	Gln	Arg	His	Glu	Gly	Asp
				725					730					735	
Leu	Lys	Glu	Lys	Glu	Arg	Leu	Val	Arg	Met	Met	Gly	Asn	Lys	Glu	Val
			740					745					750		
Tyr	GIn	Ala	Arg	Val	Val	Ser	His	Arg	Gln	Ala	Glu	Phe	Asn	Arg	Leu
		755					760					765			
Arg	Arg	Glu	Arg	Glu	Glu	Arg	11e	Ser	Arg	lle	Leu	Gln	Ser	Arg	Arg
	770					775					780				
Gln	Glu	Arg	Glu	Lys	Met	Arg	Lys	Leu	Lys	Tyr	Tyr	l.eu	Lys	Leu	Glu
785					790					795					800
Glu	Glu	Arg	Gln	Gln	Lys	Leu	Arg	Glu	Ala	Glu	Glu	Ala	Arg	Lys	Arg
				805					810					815	
Glu	Asp	Ala	Glu	Arg	Lys	Lys	Lys	Glu	Glu	Glu	Glu	Arg	Leu	Arg	Lys
			820					825					830		
Leu	Glu	Glu	lle	Ala	Glu	Lys	G1n	Arg	Gln	Arg	Glu	Arg	Glu	Leu	Glu
		835					840					845			

Glu Lys Glu Lys Gln Arg Arg Glu Ala Leu Leu Gly Arg Ala Ala Ala 855 860 Glu Pro Ala Pro Pro Ala Arg Pro Leu Glu Ser Gly Ser Ala Ala Pro 880 875 870 865 Ala Ala Ala Ala Ala Ala Ala Ala Pro Thr Pro Gly Lys Tyr Val 890 885 Pro Lys Phe Arg Arg Glu Arg Thr Glu Ser Ala Gly Ala Ala Pro Pro 905 900 Pro Glu Thr Asp Arg Trp Asn Ser Ser Ser Arg Pro Asp Gly Asp Arg 920 925 915 Trp Arg Ser Asp Asp Arg Arg Thr Ala Phe Gly Ser Gly Gly Ser 935 Arg Ser Ser Ser Thr Trp Ser Ser Ser Arg Asn Ala Arg 955 945 950

<210> 4748

<211> 177

<212> PRT

<213> Homo sapiens

<400> 4748

Met Glu Glu Asp Leu Pro Ser Lys Trp Lys Ala Lys Lys Lys Ala Gly
1 5 10 15

Leu Ala 11e Leu Val Ser Asp Lys Pro Asp Phe Lys Pro Thr Lys 11e 20 25 30

Lys Arg Asp Lys Glu Gly His Tyr IIe Val Val Lys Gly Ser Ile His
35 40 45

Gln Glu Asp Leu Thr lle Leu Asn lle Tyr Ala Pro Asn Thr Gly Ala 50 55 60

Pro Arg Ser Ile Lys Gln Val Leu Arg Asp Leu Arg Arg Asp Leu Asp .65 70 75 80

Ser His Thr 11e 11e Met Gly Asp Phe Asn Thr Pro Leu Ser 11e Leu 85 90 95

Asp Arg Ser Thr Arg Gln Lys Val 11e Lys Asp 11e Gln Asp Leu Asn 100 105 110 6926

Ser Ala Leu His Gln Ala Asp Leu Ile Asp Ile Tyr Arg Thr Leu His 120 125 His Lys Ser Arg Gln Tyr Ala Phe Phe Ser Ala Pro His Cys Thr Tyr 140 130 135 Ser Lys Ile Gly His Ile Ile Gly Ser Lys Thr Leu Leu Arg Lys Cys 155 150 Lys Arg Thr Glu Ile Thr Ala Asn Cys Leu Leu Asp Thr Val Gln Ser 170 165 Asn <210> 4749 <211> 113 <212> PRT <213> Homo sapiens <400> 4749 Met Tyr Cys Leu Lys Gln Phe Leu Pro Leu Cys Val Thr Cys Gly Leu 15 1 5 10 Ser Ile Val Pro Val Val Ile Ala Leu Ile Tyr Lys Cys Pro Leu Phe 25 Leu Ser Cys Asn Ile Phe Leu Val Thr Cys Val Thr Pro Ser Val Thr 40 45 Val His Leu Cys Ser Leu Tyr Lys His Asn Ser Ser Ser Ser Leu Cys 50 55 60 Pro Thr Pro Ser Leu Pro Phe Gly Phe Ala Cys Asn Val Phe Ile Cys 70 75 Val Arg Gln Ser Pro Thr Val Phe Leu Pro Cys Gly Ser Cys Phe Gly 85 90 Ser Leu Gln Met Ser Leu Pro Thr 11e Phe Leu Ala Ser Ser Arg Asp 100 105 110

His

<211> 202 <212> PRT <213> Homo sapiens <400> 4750 Met Thr Gly Ser Asn Ser His Ile Thr Ile Leu 11e Leu Asn Ala Asn Gly Leu Asn Ala Pro Ile Lys Arg His Arg Leu Thr Asn Trp Ile Lys Ser Gln Asp Pro Ser Val Cys Cys Ile Glu Glu Thr His Leu Met Cys Arg Asp Thr His Gly Leu Lys Ile Lys Gly Trp Arg Lys Ile Cys Gln Ala Asn Gly Lys Gln Lys Lys Glu Gly Val Ala Ile Leu Val Ser Asp Lys Thr Gly Phe Lys Pro Thr Lys Ile Lys Arg Asp Lys Glu Gly His Tyr Ile Met Val Lys Gly Ser Ile Gln Gln Glu Glu Leu Thr Val Leu Asn Ile Cys Ala Pro Asn Ile Gly Ala Pro Arg Phe Ile Lys Gln Val Leu Ser Asp Leu Gln Ser Asp Leu Asp Ser His Thr lle lle Met Gly Asp Phe Asn Thr Pro Leu Ser Thr Leu Asp Arg Ser Thr Arg Gln Lys Val Asn Lys Asp lle Gln Glu Leu Asn Thr Ala Leu His Gln Ala Asp Leu lle Asp lle Tyr Arg Thr Leu His Pro Thr Ser Thr Glu Tyr Thr Phe Phe Ser Ala Pro His Leu Phe Gln Asn

<210> 4750

<211> 512

```
<212> PRT
<213> Homo sapiens
<400> 4751
Met Gln Lys Leu Leu Arg Pro Ser Ser Val Pro Phe Leu Ser 11e Ser
                                     10
Val Thr Val Ala Thr Pro Phe Leu Ser Leu Pro Pro Lys Val Ala Asn
                                 25
                                                      30
Gln Pro Met Ser Ala Ala Ala Ala Gly Ser Ser Leu Lys Arg Gln Leu
         35
                             40
                                                 45 . . .
Ser Tyr Ser Arg Tyr Leu Ala Leu Ser Ser Thr Asn Thr Cys Ser Val
                         55
                                             60
Cys Arg Cys Leu Ser Leu Thr His Thr His Ser His Pro Lys Asn Asn
                     70
                                         75
                                                              80
65
Lys Thr Leu Leu Leu Leu Asn His Ser Arg Asn Arg Ser Leu His Ser
                 85
                                     90
Ala Ser Glu Gly Ser Phe Ile Thr Gln Pro Asp Pro Val Glu Tyr Gly
                                105
                                                     110
Ser Leu Val Asp Ser Lys Glu Lys Pro Phe Asn Ser Arg Leu Asn Arg
        115
                            120
                                                 125
Arg Gln Lys Gly Ser Thr Ser Ser Ser Pro Ala Pro Ser Asn Pro Asp
                        135
                                            140
Leu Leu Ala Ile Pro Gly Val Gly Pro Arg Asn Phe Arg Lys Leu Val
                    150
                                        155
                                                             160
145
Gln Lys Gly Ile Ala Gly Val Ala Gln Leu Lys Gln Leu Tyr Lys Asp
                165
                                    170
Lys Ser Val Asp Glu Glu Glu Leu Glu Asp Asn Ser Ser Ser Val
                                                     190
                                185
Gln Lys Lys Arg Leu Thr Phe Cys Val Glu Gly Asn Ile Ser Val Gly
                            200
                                                 205
        195
Lys Thr Thr Phe Leu Gln Arg Ile Ala Asn Glu Thr Ile Glu Leu Arg
                        215
                                            220
Asp Leu Val Glu Val Val Pro Glu Pro Ile Ser Lys Trp Gln Asp Val
225
                    230
                                        235
                                                             240
```

Gly Pro Asp His Phe Asn Ile Leu Asp Ala Phe Tyr Ala Glu Pro Gln

				245					250					255	
Arg	Tyr	Ala	Tyr	Thr	Phe	Gln	Asn	Tyr	Val	Phe	Val	Thr	Arg	Val	Met
			260					265					270		
Gln	Glu	Arg	Glu	Ser	Ser	Ala	Gly	He	Lys	Pro	Leu	۸rg	Leu	Met	Glu
		275					280					285			
Arg	Ser	Val	Phe	Ser	Asp	Arg	Met	Val	Phe	Val	Arg	Ala	Val	His	Glu
	290					295					300				
Ala	Asn	Trp	Met	Asn	Gly	Met	Glu	11e	Ser	Ile	Tyr	Asp	Ser	Trp	Phe
305					310					315					320
Asp	Pro	Val	Val	Ser	Ser	Leu	Pro	Gly	Leu	Ile	Pro	Asp	Gly	Phe	Ile
				325					330					335	
Tyr	Leu	Arg	Ala	Ser	Pro	Asp	Thr	Cys	His	Lys	Arg	Met	Met	Leu	Arg
			340					345					350		
Lys	Arg	Thr	Glu	Glu	Gly	Gly	Val	Ser	Leu	Asp	Tyr	Leu	Cys	Asp	Leu
		355					360					365			
His	Glu	Lys	His	Glu	Ser	Trp	Leu	Phe	Pro	Ser	Gln	Ser	Gly	Asn	His
	370					375					380				
Gly	Val	Leu	Ser	Val	Asn	Gln	Leu	Pro	His	His	lle	Asp	Asn	Ser	Leu
385					390					395					400
His	Pro	Asp	He	Arg	Asp	Arg	Val	Phe	Tyr	Leu	Glu	Gly	Gly	His	Met
				405					410					415	
His	Ser	Ser		Gln	Lys	Va]	Pro	Ala	Leu	Val	Leu	Asp		Glu	Pro
			420					425					430		
Asn	lle		Phe	Ser	Lys	Asp		Glu	Ala	Lys	Arg		Tyr	Ala	Arg
		435					440					445			•
Gln		Ala	Glu	Phe	Phe			Val	Lys	Lys			Glu	Val	Ser
_	450				~	455					460			В	
	Lys	Glu	G1y	Ser		GIn	Ala	Gln	Pro		Val	Leu	Leu	Pro	
465	0.1	0.1		Tr.	470	15		61		475	DI	13		C1	480
Glu	Gly	Gly	Leu		Leu	Pro	Лѕр	G1 y		Pro	Phe	Pro	Arg		Ala
,	,	C	,	485	ימ	Δ.	Cl	4.7	490	TI	C	DI.	A 1 = 4	495	C1
Leu	Lys	Ser		Asp	rhe	Arg	GIn	Ala	Ala	ihr	Ser	rne		ser	61 y
			500					505					510		

```
<211> 277
<212> PRT
<213> Homo sapiens
<400> 4752
Met Phe Val Cys Val Thr Ser Ile Thr Val Ile Ile Val Ser Lys Asp
Arg Glu Phe Cys Leu His Phe Val Met Asp Gly Ser Phe Leu Cys Ser
                                                      30
                                  25
Gln Thr Gly Lys Lys Leu Met Ala Lys Cys Arg Met Leu Ile Gln Glu
                              40
Asn Gln Glu Leu Gly Arg Gln Leu Ser Gln Gly Arg 11e Ala Gln Leu
                         55
                                              60
Glu Ala Glu Leu Ala Leu Gln Lys Lys Tyr Ser Glu Glu Leu Lys Ser
                     70
                                          75
                                                              80
 65
Ser Gln Asp Glu Leu Asn Asp Phe Ile Ile Gln Leu Asp Glu Glu Val
                                      90
                 85
Glu Gly Met Gln Ser Thr Ile Leu Val Leu Gln Gln Gln Leu Lys Glu
                                                     110
                                 105
Thr Arg Gln Gln Leu Ala Gln Tyr Gln Gln Gln Gln Ser Gln Ala Ser
                                                 125
        115
                             120
Ala Pro Ser Thr Ser Arg Thr Thr Ala Ser Glu Pro Val Glu Gln Ser
                        135
                                             140
Glu Ala Thr Ser Lys Asp Cys Ser Arg Leu Thr Asn Gly Pro Ser Asn
                                                              160
                    150
                                         155
145
Gly Ser Ser Ser Arg Gln Arg Thr Ser Gly Ser Gly Phe His Arg Glu
                                     170
                165
Gly Asn Thr Thr Glu Asp Asp Phe Pro Ser Ser Pro Gly Asn Gly Asn
                                                     190
                                 185
Lys Ser Ser Asn Ser Ser Glu Glu Arg Thr Gly Arg Gly Gly Ser Gly
                             200
                                                 205
        195
Tyr Val Asn Gln Leu Ser Ala Gly Tyr Glu Ser Val Asp Ser Pro Thr
                        215
                                             220
Gly Ser Glu Asn Ser Leu Thr His Gln Ser Asn Asp Thr Asp Ser Ser
                                                              240
225
                    230
                                         235
```

His Asp Pro Gln Glu Glu Lys Ala Val Ser Gly Lys Gly Asn Arg Thr

Val Gly Ser Arg His Val Gln Asn Gly Leu Asp Ser Ser Val Asn Val Gln Gly Ser Val Leu <210> 4753 ⟨211⟩ 528 <212> PRT <213> Homo sapiens <400> 4753 Met Ala Val Gly Gln Leu Val Pro Ser Asn Leu Phe Leu Asn His Ser Ala Ala Thr Thr Leu Ile Pro Thr Ser Thr Pro Pro Arg His Arg His Leu Leu Cys Ile Ser Ser Ala Asn His Thr Thr Ala Thr Asp Asn Asp Ser Pro Phe Pro Ser Phe Gly Arg Val Lys Thr Leu Leu Val His Arg Arg Arg Lys Asp Gln Ser His Arg Arg Ala Val Gln Leu Glu Asp Asp Asn Asp Asp Ile Ala Pro Arg Pro Arg Arg Ser Gln Ser Arg Ser Arg Gly Gly Glu Arg Trp Asp Met Ile Pro Asn Tyr Thr Pro Gln Ser Lys Ser Ala Thr Asp Thr Lys Phe Phe Ser Leu Lys Ser Phe Lys Glu lle Gly Cys Ser Glu Tyr Met Ile Glu Ser Leu Gln Lys Leu Leu Leu Ser Arg Pro Ser His Val Gln Ala Met Ala Phe Ala Pro Val Ile Ser Gly Lys Thr Cys Val Ile Ala Asp Gln Ser Gly Ser Gly Lys Thr Leu

Ala Tyr Leu Ala Pro Ile Ile Gln Leu Leu Arg Leu Glu Glu Leu Glu

			180					185					190		
G1 y	Arg	Ser	Ser	Lys	Ser	Ser	Ser	Gln	Ala	Pro	Arg	Val	Leu	Val	Leu
		195					200					205			
Ala	Pro	Thr	Ala	Glu	Leu	Ala	Ser	Gln	Val	Leu	Asp	Asn	Cys	Arg	Ser
	210					215					220				
Leu	Ser	Lys	Ser	Gly	Val	Pro	Phe	Lys	Ser	Met	Val	Val	Thr	Gly	Gly
225					230					235					240
Phe	Arg	Gln	Lys	Thr	Gln	Leu	Glu	Asn	Leu	Gln	Gln	Gly	Val	Asp	Val
				245					250					255	
Leu	Ile	Ala	Thr	Pro	Gly	Arg	Phe	Leu	Phe	Leu	Ile	His	Glu	G1 y	Phe
			260					265					270		
Leu	Gln	Leu	Thr	Asn	Leu	Arg	Cys	Ala	Ile	Leu	Asp	Glu	Val	Asp	lle
		275					280					285			
Leu	Phe	Gly	Asp	Glu	Asp	Phe	Glu	Val	Ala	Leu	Gln	Ser	Leu	11e	Asn
	290					295					300				
Ser	Ser	Pro	Val	Asp	Thr	Gln	Tyr	Leu	Phe	Val	Thr	Ala	Thr	Leu	Pro
305					310					315					320
Lys	Asn	Val	Tyr	Thr	Lys	Leu	Val	Glu	Val	Phe	Pro	Asp	Cys	Glu	Met
				325					330					335	
He	Met	Gly	Pro	Gly	Met	His	Arg	Ile	Ser	Ser	Arg	Leu	Gln	Glu	Ile
			340					345					350		
He	Val	Asp	Cys	Ser	Gly	Glu	Asp	Gly	Gln	Glu	Lys	Thr	Pro	Asp	Thr
		355					360					365			
Ala	Phe	Leu	Asn	Lys	Lys	Thr	Ala	Leu	Leu	Gln	Leu	Val	Glu	Glu	Asn
	370					375					380				
Pro	Val	Pro	Arg	Thr	lle	Val	Phe	Cys	Asn	Lys	lle	Glu	Thr	Cys	Arg
385					390					395					400
Lys	Val	Glu	Asn	Leu	Leu	Lys	Arg	Phe	Asp	Arg	Lys	Gly	Asn	His	Val
				405					410					415	
Gln	Val	Leu	Pro	Phe	His	Ala	Ala	Met	Thr	Gln	Glu	Ser	Arg	Leu	Ala
			420					425					430		
Ser	Met	Glu	Glu	Phe	Th:r	Arg	Ser	Pro	Ser	Lys	Gly	Val	Ser	Gln	Phe
		435					440					445			
Met	Val	Cys	Thr	Asp	Arg	Ala	Ser	Arg	Gly	He	Asp	Phe	Thr	Arg	Val
	450					455					460				
Asp	His	Val	He	Leu	Phe	Asp	Phe	Pro	Arg	Asp	Pro	Ser	Glu	Tyr	Val

Arg Arg Val Gly Arg Thr Ala Arg Gly Ala Lys Gly Val Gly Lys Ala Phe Ile Phe Val Val Gly Lys Gln Val Ser Leu Ala Arg Lys Ile Met Glu Arg Asn Gln Lys Gly His Pro Leu His Asp Val Pro Ser Ala Tyr <210> 4754 <211> 238 <212> PRT <213> Homo sapiens <400> 4754 Met Cys Gly Leu Ser Phe Met Leu Phe Arg Ala Cys Leu Lys Met Asp Leu Gly Val Phe Ile Cys Val Phe Ile Ser Phe Ser Pro Leu Phe Lys Ser Thr Ala Asp Cys Pro Asp Ala Val Pro Ser Ser Ala Glu Thr Gly Gly Thr Asn Tyr Leu Ala Pro Gly Gly Leu Ser Asp Ser Gln Leu Leu Leu Glu Pro Gly Asp Arg Ser His Trp Cys Val Val Ala Tyr Trp Glu Glu Lys Thr Arg Val Gly Arg Leu Tyr Cys Val Gln Glu Pro Ser Leu Asp lle Phe Tyr Asp Leu Pro Gln Gly Asn Gly Phe Cys Leu Gly Gln Leu Asn Ser Asp Asn Lys Ser Gln Leu Val Gln Lys Val Arg Ser Lys lle Gly Cys Gly lle Gln Leu Thr Arg Glu Val Asp Gly Val Trp Val Tyr Asn Arg Ser Ser Tyr Pro Ile Phe Ile Lys Ser Ala Thr Leu Asp

Asn Pro Asp Ser Arg Thr Leu Leu Val His Lys Val Phe Pro Gly Phe Ser Ile Lys Ala Phe Asp Tyr Glu Lys Ala Tyr Ser Leu Gln Arg. Pro Asn Asp His Glu Phe Met Gln Gln Pro Trp Thr Gly Phe Thr Val Gln lle Ser Phe Val Lys Gly Trp Gly Gln Cys Tyr Thr Arg Gln Phe lle Ser Ser Cys Pro Cys Trp Leu Glu Val Ile Phe Asn Ser Arg

<210> 4755

<211> 307

<212> PRT

<213> Homo sapiens

<400> 4755

Met Gln Ala Leu Ser Thr Val Pro Leu Asp Trp Val Thr Val Pro Lys Leu Gln Glu Cys Gly Ala Arg Pro Ala Met Glu Lys Pro Thr Arg Val Val Gly Gly Phe Gly Ala Ala Ser Gly Glu Val Pro Trp Gln Val Ser Leu Lys Glu Gly Ser Arg His Phe Cys Gly Ala Thr Val Val Gly Asp Arg Trp Leu Leu Ser Ala Ala His Cys Phe Asn His Thr Lys Val Glu Gln Val Arg Ala His Leu Gly Thr Ala Ser Leu Leu Gly Leu Gly Gly Ser Pro Val Lys lle Gly Leu Arg Arg Val Val Leu His Pro Leu Tyr Asn Pro Gly 11e Leu Asp Phe Asp Leu Ala Val Leu Glu Leu Ala Ser Pro Leu Ala Phe Asn Lys Tyr lle Gln Pro Val Cys Leu Pro Leu Ala

Ile Gln Lys Phe Pro Val Gly Arg Lys Cys Met Ile Ser Gly Trp Gly Asn Thr Gln Glu Gly Asn Ala Thr Lys Pro Glu Leu Leu Gln Lys Ala Ser Val Gly Ile 11e Asp Gln Lys Pro Cys Ser Val Leu Tyr Asn Phe Ser Leu Thr Asp Arg Met Ile Cys Ala Gly Phe Leu Glu Gly Lys Val Asp Ser Cys Gln Gly Asp Ser Gly Gly Pro Leu Ala Cys Glu Glu Ala Pro Gly Val Phe Tyr Leu Ala Gly Ile Val Ser Trp Gly Ile Gly Cys Ala Gln Val Lys Lys Pro Gly Val Tyr Thr Arg lle Thr Arg Leu Lys Gly Trp Ile Leu Glu Ile Met Ser Ser Gln Pro Leu Pro Met Ser Pro Pro Ser Thr Thr Arg Met Leu Ala Thr Thr Ser Pro Arg Thr Thr Ala Gly Leu Thr Val Pro Gly Ala Thr Pro Ser Arg Pro Thr Pro Gly Leu Pro Ala Gly

<210> 4756

<211> 583

<212> PRT

<213> Homo sapiens

<400> 4756

 Met Ser Val 11e Val Lys Thr Gln Glu Gly Asp 11e Leu Leu Phe Cys

 1
 5
 10
 15

 Lys Gly Ala Asp Ser Ala Val Phe Pro Arg Val Gln Asn His Glu 11e
 20
 25
 30

 Glu Leu Thr Lys Val His Val Glu Arg Asn Ala Met Asp Gly Tyr Arg
 35
 40
 45

Thr	Leu	Cys	Val	Ala	Phe	Lys	Glu	lle	Ala	Pro	Asp	Asp	Tyr	Glu	Arg
	50					55					60				
He	Asn	Arg	Gln	Leu	He	Glu	Ala	Lys	Met	Ala	Leu	Gln	Asp	Arg	Glu
65					70					75					80
Glu	Lys	Met	Glu	Lys	Val	Phe	Asp	Λsp	Пе	Glu	Thr	Asn	Met	Asn	Leu
				85					90					95	
He	Gly	Ala	Thr	Ala	Val	Glu	Asp	Lys	Leu	Gln	Asp	Gln	Ala	Ala	Glu
			100					105					110		
Thr	He	Glu	Ala	Leu	His	Ala	Ala	G1 y	Leu	Lys	Val	Trp	Val	Leu	Thr
		115					120					125			
Gly	Asp	Lys	Met	Glu	Thr	Ala	Lys	Ser	Thr	Cys	Tyr	Ala	Cys	Arg	Leu
	130					135					140				
Phe	Gln	Thr	Asn	Thr	Glu	Leu	Leu	Glu	Leu	Thr	Thr	Lys	Thr	He	Glu
145					150					155					160
Glu	Ser	Glu	Arg	Lys	Glu	Asp	Arg	Leu	His	Glu	Leu	Leu	lle	Glu	Tyr
				165					170					175	
Arg	Lys	Lys		Leu	His	Glu	Phe		Lys	Ser	Thr	Arg		Phe	Lys
			180					185					190		
Lys	Ala		Thr	Glu	His	Gln		Tyr	Gly	Leu	Ile		Asp	Gly	Ser
	_	195			_		200	_			_	205			
Thr		Ser	Leu	He	Leu	Asn	Ser	Ser	GIn	Asp	Ser	Ser	Ser	Asn	Asn
m.															
	210	6		D.	•	215		0			220		. 1	., .	•
225		Ser	lle	Phe		215 Gln	He	Cys	Met			Thr	Ala	Va1	
Cys	Lys				230	Gln				235	Cys				240
	Lys			Ala	230				Ala	235	Cys			Met	240
Luc	Lys Cys	Arg	Met	Ala 245	230 Pro	G1n Leu	Gln	Lys	Ala 250	235 Gln	Cys Ile	Val	Arg	Met 255	240 Val
Lys	Lys Cys	Arg	Met Lys	Ala 245	230 Pro	Gln	Gln	Lys Thr	Ala 250	235 Gln	Cys Ile	Val	Arg Asp	Met 255	240 Val
	Lys Cys Asn	Arg Leu	Met Lys 260	Ala 245 Gly	230 Pro Ser	Gln Leu Pro	Gln lle	Lys Thr 265	Ala 250 Leu	235 Gln Ser	Cys lle	Val Gly	Arg Asp 270	Met 255 Gly	240 Val Ala
	Lys Cys Asn	Arg Leu Val	Met Lys 260	Ala 245 Gly	230 Pro Ser	G1n Leu	Gln lle Glu	Lys Thr 265	Ala 250 Leu	235 Gln Ser	Cys lle	Val Gly lle	Arg Asp 270	Met 255 Gly	240 Val Ala
Asn	Lys Cys Asn Asp	Arg Leu Val 275	Met Lys 260 Ser	Ala 245 Gly Met	230 Pro Ser 11e	Gln Leu Pro Leu	Gln 11e Glu 280	Lys Thr 265 Ser	Ala 250 Leu His	235 Gln Ser Val	Cys Ile Ile Gly	Val Gly 11e 285	Arg Asp 270 Gly	Met 255 Gly Ile	240 Val Ala Lys
Asn	Lys Cys Asn Asp Lys	Arg Leu Val 275	Met Lys 260 Ser	Ala 245 Gly Met	230 Pro Ser 11e	Gln Leu Pro Leu Ala	Gln 11e Glu 280	Lys Thr 265 Ser	Ala 250 Leu His	235 Gln Ser Val	Cys Ile Ile Gly Asp	Val Gly 11e 285	Arg Asp 270 Gly	Met 255 Gly Ile	240 Val Ala Lys
Asn Gly	Lys Cys Asn Asp Lys 290	Arg Leu Val 275 Glu	Met Lys 260 Ser Gly	Ala 245 Gly Met	230 Pro Ser 11e Gln	Gln Leu Pro Leu Ala 295	Gln 11e Glu 280 Ala	Lys Thr 265 Ser Arg	Ala 250 Leu His	235 Gln Ser Val	Cys Ile Ile Gly Asp 300	Val Gly 11e 285 Tyr	Arg Asp 270 Gly Ser	Met 255 Gly Ile Val	240 Val Ala Lys Pro
Asn Gly Lys	Lys Cys Asn Asp Lys 290	Arg Leu Val 275 Glu	Met Lys 260 Ser Gly	Ala 245 Gly Met	230 Pro Ser Ile Gln Lys	Gln Leu Pro Leu Ala	Gln 11e Glu 280 Ala	Lys Thr 265 Ser Arg	Ala 250 Leu His	235 Gln Ser Val Ser	Cys Ile Ile Gly Asp 300	Val Gly 11e 285 Tyr	Arg Asp 270 Gly Ser	Met 255 Gly Ile Val	240 Val Ala Lys Pro
Asn Gly Lys 305	Lys Cys Asn Asp Lys 290 Phe	Arg Leu Val 275 Glu Lys	Met Lys 260 Ser Gly	Ala 245 Gly Met Arg	230 Pro Ser 11e Gln Lys 310	Gln Leu Pro Leu Ala 295	Gln 11e Glu 280 Ala Leu	Lys Thr 265 Ser Arg	Ala 250 Leu His Asn	235 Gln Ser Val Ser Ala 315	Cys Ile Ile Gly Asp 300 His	Val Gly 11e 285 Tyr	Arg Asp 270 Gly Ser	Met 255 Gly Ile Val	240 Val Ala Lys Pro Tyr 320

Cys	Phe	lle	Leu	Pro	Gln	Phe	Leu	Tyr	Gln	Phe	Phe	Cys	Gly	Phe	Ser
			340					345					350		
Gln	Gln	Pro	Leu	Tyr	Asp	Ala	Ala	Tyr	Leu	Thr	Met	Tyr	Asn	Пе	Cys
		355					360					365			
Phe	Thr	Ser	Leu	Pro	Пe	Leu	Ala	Tyr	Ser	Leu	Leu	Glu	Gln	His	He
	370					375					380				
Asn	Пe	Asp	Thr	Leu	Thr	Ser	Asp	Pro	Arg	Leu	Tyr	Met	Lys	lle	Ser
385					390					395					400
Gly	Asn	Ala	Met	Leu	Gln	Leu	Gly	Pro	Phe	Leu	Tyr	Trp	Thr	Phe	Leu
				405					410					415	
Ala	Ala	Phe	Glu	Gly	Thr	Val	Phe	Phe	Phe	Gly	Thr	Tyr	Phe	Leu	Phe
			420					425					430		
Gln	Thr	Ala	Ser	Leu	Glu	Glu	Asn	G1 y	Lys	Val	Tyr	Gly	Asn	Trp	Thr
		435					440					445			
Phe	Gly	Thr	lle	Val	Phe	Thr	Val	Leu	Va]	Phe	Thr	Va]	Thr	Leu	Lys
	450					455					460				
Leu	Ala	Leu	Asp	Thr	Arg	Phe	Trp	Thr	Trp	lle	Asn	His	Phe	Val	lle
465					470					475					480
Trp	Gly	Ser	Leu	Ala	Phe	Tyr	Val	Phe	Phe	Ser	Phe	Phe	Trp	Gly	Gly
				485					490					495	
He	lle	Trp	Pro	Phe	Leu	Lys	Gln	Gln	Arg	Met	Tyr	Phe	Val	Phe	Ala
			500					505					510		
Gln	Met	Leu	Ser	Ser	Val	Ser	Thr	Trp	Leu	Ala	He	lle	Leu	Leu	He
		515					520					525			
Phe	lle	Ser	Leu	Phe	Pro		lle	Leu	Leu	He	Val	Leu	Lys	Asn	Val
	530					535					540				
Arg	Arg	Arg	Ser	Ala	Arg	Arg	Asn	Leu	Ser		Arg	Arg	Ala	Ser	
545					550					555					560
Ser	Leu	Ser	Ala		Pro	Ser	Val	Arg		Leu	Leu	Leu	Arg		Phe
				565					570					575	
Ser	Asp	Glu		Asn	Val	Leu									
			580												

<210> 4757 <211> 171 <212> PRT

<213> Homo sapiens

<400> 4757

Met His Leu Trp Arg Tyr Pro Ser Leu Ser Ile His Gly 11e Glu Gly

5 10 1

Ala Phe Asp Glu Pro Gly Thr Lys Thr Val Ile Pro Gly Arg Val Ile

20 25 30

Gly Lys Phe Ser Ile Arg Leu Val Pro His Met Asn Val Ser Ala Val
35 40 45

Glu Lys Gln Val Thr Arg His Leu Glu Asp Val Phe Ser Lys Arg Asn 50 55 60

Ser Ser Asn Lys Met Val Val Ser Met Thr Leu Gly Leu His Pro Trp
65 70 75 80

11e Ala Asn 11e Asp Asp Thr Gln Tyr Leu Ala Ala Lys Arg Ala I1e 85 90 95

Arg Thr Val Phe Gly Thr Glu Pro Asp Met Ile Arg Asp Gly Ser Thr
100 105 110

Ile Pro Ile Ala Lys Met Phe Gln Glu Ile Val His Lys Ser Val Val

115 120 125

Leu 11e Pro Leu Gly Ala Val Asp Asp Gly Glu His Ser Gln Asn Glu 130 135 140

Lys 11e Asn Arg Ser Ala Asp Ala Cys Ala Met Cys Leu Ser Leu Leu 145 150 155 160

Pro Leu Leu His Thr Pro Gly Ser Thr Arg Gly

165 170

<210> 4758

<211> 446

<212> PRT

<213> Homo sapiens

<400> 4758

Met Leu Gly Gly Val Val Glu Leu Ala Ala Ala Ser Ser Phe Pro Leu

1				5					10					15	
Leu	Glu	Gln	Phe	Ala	Gly	Asp	Gly	Phe	His	Ala	Asp	Gly	Asp	Asp	Val
			20					25					30		
Leu	Thr	Glu	Leu	Cys	Ala	Pro	Asp	Gly	Ala	Pro	Pro	Gly	Val	Val	Pro
		35					40					45			
Val	Leu	Ser	Ala	His	Ser	Pro	Ser	Leu	Gly	Ser	Glu	Tyr	Phe	He	Arg
	50					55					60				
Leu	Glu	Glu	Ala	Ala	Pro	Ala	Thr	Gly	His	Asp	Pro	Asp	Cys	Ala	Gly
65					70					75					80
Cys	Ala	Pro	Ser	Pro	Pro	Ala	He	Ala	Asp	Gln	Asp	Asp	Asp	Ser	Asp
				85					90					95	
Gly	Ser	Thr	Ala	Ala	Ser	Leu	Ala	Met	G]u	Pro	Leu	Leu	G1 y	His	Gly
			100					105					110		
Pro	Pro	Val	Asp	Va]	Pro	Trp	Gly	Arg	Gly	Asp	His	Tyr	Pro	Arg	Arg
		115					120					125			
Ser	Leu	Ala	Arg	Asp	Pro	Leu	Cys	Pro	Ser	Arg	Ser	P.ro	Ser	Pro	Ser
	130					135					140				
Ala	Gly	Pro	Leu	Ser	Leu	Ala	Glu	Gly	Gly	Ala	Glu	Asp	Ala	Asp	Trp
145					150					155					160
Gly	Val	Ala	Ala	Phe	Cys	Pro	Ala	Phe	Phe	Glu	Asp	Pro	Leu	Gly	Thr
				165					170					175	
Ser	Pro	Leu	Gly	Ser	Ser	G1 y	Ala	Pro	Pro	Leu	Pro	Leu	Thr	Gly	Glu
			180					185					190		
Asp	Glu	Leu	Glu	Glu	Val	Gly	Ala	Arg	Arg	Ala	Ala	Gln	Arg	Gly	His
		195					200					205			
Trp	Arg	Ser	Asn	Val	Ser	Ala	Asn	Asn	Asn	Ser	Gly	Ser	Arg	Cys	Pro
	210					215					220				
	Ser	Trp	Asp	Pro	Val	Ser	Ala	Gly	Gly	His	Ala	Glu	Gly	Cys	Pro
225					230					235					240
Ser	Pro	Lys	Gln		Pro	Arg	Ala	Ser		Glu	Pro	Gly	Tyr		G] y
				245					250					255	
Glu	Pro	Leu		Gly	Leu	G]n	Ala		Ser	Ala	Gln	Glu		Gly	Cys
			260					265					270		
Cys	Pro		Leu	Pro	His	Leu		Ser	Ala	Gln	Gly		Ala	Pro	Ala
_	_	275					280					285			
Pro	Cys	Leu	Val	Thr	Pro	Ser	Trp	Thr	Glu	Thr	Ala	Ser	Ser	GIy	Gly

290	295	300	
Asp His Pro Gln A	Na Glu Pro Lys	Leu Ala Thr Glu Ala	Glu Gly Thr
305	310	315	320
Thr Gly Pro Arg L	leu Pro Leu Pro	Ser Val Pro Ser Pro	Ser Gln Glu
3	325	330	335
Gly Ala Pro Leu P	Pro Ser Glu Glu	Ala Ser Ala Pro Asp	Ala Pro Asp
340		345	350
Ala Leu Pro Asp S	Ser Prọ Thr Pro	Ala Thr Gly Gly Glu	Val Ser Ala
355	360	365	
Ile Lys Leu Ala S	Ser Ala Leu Asn	Gly Ser Ser Ser Ser	Pro Glu Val
370	375	380	
Glu Ala Pro Ser S	Ser Glu Asp Glu	Asp Thr Ala Glu Ala	Thr Ser Gly
385	390	395	400
lle Phe Thr Asp T	Thr Ser Ser Asp	Gly Leu Gln Ala Arg	Arg Pro Asp
4	105	410	415
Val Val Pro Ala P	Phe Arg Ser Leu	Gln Lys Gln Trp Pro	Gln Arg Glu
420		425	430
Glu Ser Leu Pro A	Arg Leu Cys Leu	Leu Leu Arg Pro Arg	Gly
435	440	445	

<210> 4759

<211> 106

<212> PRT

<213> Homo sapiens

<400> 4759

Gln Ala Leu Ile Thr Leu Ser Ser Cys Ser Leu Gly Lys Cys Thr Phe

65 70 75 Tyr Thr Asn Val Arg Gln Phe Pro Gln Leu Pro Leu Leu Asp 11e Lys Glu Phe Thr Phe Leu Thr Ile Phe Gln Ser 100 105 <210> 4760 <211> 107 <212> PRT <213> Homo sapiens <400> 4760 Met Tyr Leu Leu Ile Met Glu Phe Met Ser Phe Pro Ser Phe Ala Trp 1 5 10 15 Pro Arg Phe Phe Phe Ser Leu Lys Phe Leu Leu Arg Cys Tyr Leu Gly 25 Cys Leu His Tyr Lys Gln Ala Ala Leu Pro Pro Leu Phe Val Val Thr 45 40 Gly Glu Lys Pro Tyr Lys Cys Thr Trp Glu Gly Cys Asp Trp Arg Phe 50 55 60 Ala Arg Ser Asp Glu Leu Thr Arg His Tyr Arg Lys His Thr Gly Ala 70 75 Lys Pro Phe Gln Cys Gly Val Cys Asn Arg Ser Phe Ser Arg Ser Asp 85 90 95 His Leu Ala Leu His Met Lys Arg His Gln Asn 100 105

<210> 4761

<211> 667

<212> PRT

<213> Homo sapiens

<400> 4761

Met Glu Asp Leu Ser Ser Pro Asp Ser Thr Leu Leu Gln Gly Gly His

1				5					10					15	
Asn	Leu	Leu	Ser	Ser	Ala	Ser	Phe	Gln	Glu	Ala	Val	Thr	Phe	Lys	Asp
			20					25					30		
Val	Ile	Val	Asp	Phe	Thr	Gln	Glu	Glu	Trp	Lys	Gln	Leu	Asp	Pro	Gly
		35					40					45			
Gln	Arg	Asp	Leu	Phe	Arg	Asp	Val	Thr	Leu	Glu	Asn	Tyr	Thr	His	Leu
	50					55					60				
Val	Ser	lle	Gly	Leu	Gln	Val	Ser	Lys	Pro	Asp	Val	lle	Ser	Gln	Leu
65					70					75					80
Glu	Gln	Gly	Thr	Glu	Pro	Trp	lle	Met	Glu	Pro	Ser	Ile	Pro	Val	Gly
				85					90					95	
Thr	Cys	Ala	Asp	Trp	Glu	Thr	Arg	Leu	Glu	Asn	Ser	Val	Ser	Ala	Pro
			100					105					110		
Glu	Pro	Asp	Џе	Ser	Glu	Glu	Glu	Leu	Ser	Pro	G]u	Val	lle	Val	Glu
		115					120					125			
Lys	His	Lys	Arg	Asp	Asp	Ser	Trp	Ser	Ser	Asn	Leu	Leu	Glu	Ser	Trp
	130					135					140				
Glu	Tyr	Glu	Gly	Ser	Leu	Glu	Arg	Gln	Gln	Ala	Asn	Gln	Gln	Thr	Leu
145					150					155					160
Pro	Lys	Glu	He	Lys	Val	Thr	Glu	Lys	Thr	lle	Pro	Ser	Trp	G1u	Lys
				165					170					175	
G1 y	Pro	Val		Asn	G1u	Phe	Gly		Ser	Val	Asn	Val	Ser	Ser	Asn
			180					185					190		
Leu	Val		Gln	Glu	Pro	Ser		Glu	Glu	Thr	Ser		Lys	Arg	Ser
		195		_			200					205			
He		GIn	Asn	Ser	Asn	Pro	Val	Lys	Lys	Glu		Ser	Cys	Lys	Cys
	210					215		_	~		220		~ ~		
	Glu	Cys	Gly	Lys		Phe	Ser	Tyr	Cys		Ala	Leu	He	Arg	
225		m)	,	TO I	230	0.1			Tr.	235			0.1	0	240
GIn	Arg	lhr	HIS		Gly	Glu	Lys	Pro		Lys	Cys	Asn	Glu		Glu
,	4.7	DI	C	245	C	C1			250			0.1		255	
Lys	Ala	rne		Arg	5er	Glu	Asn		не	Asn	H1S	GIN		116	H1S
TL:	C1	Λ	260	D	т	1	C	265	C I	C	C1	1	270	D⊩	т 1
ınr	oly	275	Lys	1.0	ıyr	Lys	280	лsр	GIN	Cys	01 y	285	GIY	rne	116

Glu	Gly	Pro	Ser	Leu	Thr		His	Gln	Arg	Ile		Thr	Gly	Glu	Lys
	290					295					300				
Pro	Tyr	Lys	Cys	Asp	Glu	Cys	Gly	Lys	Ala	Phe	Ser	G1n	Arg	Thr	His
305					310					315					320
Leu	Val	Gln	His	Gln	Arg	He	His	Thr	Gly	Glu	Lys	Pro	Tyr	Thr	Cys
				325					330					335	
Asn	Glu	Cys	Gly	Lys	Ala	Phe	Ser	Gln	Arg	Gly	His	Phe	Met	Glu	llis
			340					345					350		
Gln	Lys	Ile	His	Thr	Gly	Glu	Lys	Pro	Phe	Lys	Cys	Asp	Glu	Cys	Asp
		355					360					365			
Lys	Thr	Phe	Thr	Arg	Ser	Thr	His	Leu	Thr	Gln	His	Gln	Lys	He	His
	370					375					380				
Thr	Gly	Glu	Lys	Thr	Tyr	Lys	Cys	Asn	Glu	Cys	Gly	Lys	Ala	Phe	Asn
385					390					395					400
Gly	Pro	Ser	Thr	Phe	lle	Arg	His	llis	Met	Ile	His	Thr	G1 y	Glu	Lys
				405					410					415	
Pro	Tyr	Glu	Cys	Asn	Glu	Cys	Gly	Lys	Ala	Phe	Ser	Gln	His	Ser	Asn
			420					425					430		
Leu	Thr	Gln	His	Gln	Lys	Thr	His	Thr	Gln	Glu	Lys	Ala	Tyr	Glu	Cys
		435					440					445			
Lys	Glu	Cys	Gly	Lys	Ala	Phe	lle	Arg	Ser	Ser	Ser	Leu	Ala	Lys	His
	450					455					460				
Glu	Arg	Ile	His	Thr	Gly	G] u	Lys	Pro	Tyr	Gln	Cys	His	Glu	Cys	Gly
465					470					475					480
Lys	Thr	Phe	Ser	Tyr	Gly	Ser	Ser	Leu	Ile	Gln	His	Arg	Lys	He	His
				485					490					495	
Thr	Gly	Glu	Arg	Pro	Tyr	Lys	Cys	Asn	Glu	Cys	Gly	Arg	Ala	Phe	Asn
			500					505					510		
Gln	Asn	lle	His	Leu	Thr	Gln	llis	Lys	Arg	Ile	His	Thr	Gly	Ala	Lys
		515					520					525			
Pro	Tyr	Glu	Cys	Ala	Glu	Cys	Gly	Lys	Ala	Phe	Arg	llis	Cys	Ser	Ser
	530					535					540				
Leu	Ala	Gln	His	Gln	Lys	Thr	His	Thr	Glu	Glu	Lys	Pro	Tyr	Gln	Cys
545					550					555					560
Asn	Lys	Cys	Glu	Lys	Thr	Phe	Ser	Gln	Ser	Ser	His	Leu	Thr	Gln	His
				565					570					575	

Gln Arg Ile His Thr Gly Glu Lys Pro Tyr Lys Cys Asn Glu Cys Asp Lys Ala Phe Ser Arg Ser Thr His Leu Thr Glu His Gln Asn Thr His Thr Gly Glu Lys Pro Tyr Asn Cys Asn Glu Cys Arg Lys Thr Phe Ser Gln Ser Thr Tyr Leu lle Gln His Gln Arg Ile His Ser Gly Glu Lys Pro Phe Gly Cys Asn Asp Cys Gly Lys Ser Phe Arg Tyr Arg Ser Ala Leu Asn Lys His Gln Arg Leu His Pro Gly Ile

<210> 4762

<211> 460

<212> PRT

<213> Homo sapiens

<400> 4762

Met Glu Val Leu Glu Ser Gly Glu Gln Gly Val Leu Gln Trp Asp Arg Lys Leu Ser Glu Leu Ser Glu Pro Gly Asp Gly Glu Ala Leu Met Tyr His Thr His Phe Ser Glu Leu Leu Asp Glu Phe Ser Gln Asn Val Leu Gly Gln Leu Leu Asn Asp Pro Phe Leu Ser Glu Lys Ser Val Ser Met Glu Val Glu Pro Ser Pro Thr Ser Pro Ala Pro Leu lle Gln Ala Glu His Ser Tyr Ser Leu Cys Glu Glu Pro Arg Ala Gln Ser Pro Phe Thr His lle Thr Thr Ser Asp Ser Phe Asn Asp Asp Glu Val Glu Ser Glu Lys Trp Tyr Leu Ser Thr Asp Phe Pro Ser Thr Ser Ile Lys Thr Glu

Pro	Val	Thr	Asp	Glu	Pro	Pro	Pro	Gly	Leu	Val	Pro	Ser	Val	Thr	Leu
	130					135					140				
Thr	lle	Thr	Ala	11e	Ser	Thr	Pro	Leu	Glu	Lys	Glu	Glu	Pro	Pro	Leu
145					150					155					160
Glu	Met	Asn	Thr	Gly	Val	Asp	Ser	Ser	Cys	Gln	Thr	He	He	Pro	Lys
				165					170					175	
Ile	Lys	Leu	Glu	Pro	His	Glu	Val	Asp	Gln	Phe	Leu	Asn	Phe	Ser	Pro
			180					185					190		
Lys	Glu	Ala	Pro	Val	Asp	His	Leu	His	Leu	Pro	Pro	Thr	Pro	Pro	Ser
		195					200					205			
Ser	His	Gly	Ser	Asp	Ser	Glu	Gly	Ser	Leu	Ser	Pro	Asn	Pro	Arg	Leu
	210					215					220				
His	Pro	Phe	Ser	Leu	Pro	Gln	Thr	His	Ser	Pro	Ser	Arg	Ala	Ala	Pro
225					230					235					240
Arg	Ala	Pro	Ser	Ala	Leu	Ser	Ser	Ser	Pro	Leu	Leu	Thr	Ala	Pro	His
				245					250					255	
Lys	Leu	Gln	G1 y	Ser	Gly	Pro	Leu	Val	Leu	Thr	Glu	Glu	Glu	Lys	Arg
			260					265					270		
Thr	Leu	lle	Ala	Glu	Gly	Tyr	Pro	lle	Pro	Thr	Lys	Leu	Pro	Leu	Ser
		275					280					285			
Lys	Ser	Glu	Glu	Lys	Ala	Leu	Lys	Lys	lle	Arg	Arg	Lys	Ile	Lys	Asn
	290					295					300				
Lys	He	Ser	Ala	Gln	Glu	Ser	Arg	Arg	Lys	Lys	Lys	Glu	Tyr	Met	Asp
305					310					315					320
Ser	Leu	Glu	Lys	Lys	Val	Glu	Ser	Cys	Ser	Thr	Glu	Asn	Leu	Glu	Leu
				325					330					335	
Arg	Lys	Lys	Va]	Glu	Val	Leu	Glu	Asn	Thr	Asn	Arg	Thr	Leu	Leu	GIn
			340					345					350		
Gln	Leu	Gln	Lys	Leu	Gln	Thr	Leu	Val	Met	Gly	Lys	Val	Ser	Arg	Thr
		355					360					365			
Cys	Lys	Leu	Ala	Gly	Thr	Gln	Thr	Gly	Thr	Cys	Leu	Met	Val	Val	Va]
	370					375					380				
Leu	Cys	Phe	Ala	Val	Ala	Phe	Gly	Ser	Phe	Phe	Gln	Gly	Tyr	Gly	Pro
385					390					395					400
Tyr	Pro	Ser	Ala	Thr	Lys	Met	Ala	Leu	Pro	Ser	Gln	His	Ser	Leu	Gln
				405					410					415	

Glu Pro Tyr Thr Ala Ser Val Gly Lys Thr Ala Cys Gly Lys Leu Gly Arg Val Leu Phe Tyr Phe Pro Arg Ala Gly Phe Leu Ser Leu Pro Lys Gly Ile Phe Cys Glu Ser Pro Met Phe Lys Lys Trp

<210> 4763

<211> 443

<212> PRT

<213> Homo sapiens <400> 4763 Met Glu His Val Gln Gly Arg lle Phe Arg Asp Leu Thr Ile Pro Gly Leu Ser Pro Ala Glu Arg Ser Ala lle Tyr Val Ala Thr Val Glu Thr Leu Ala Gln Leu His Ser Leu Asn Ile Gln Ser Leu Gln Leu Glu Gly Tyr Gly lle Gly Ala Gly Tyr Cys Lys Arg Gln Val Ser Thr Trp Thr Lys Gln Tyr Gln Ala Ala Ala His Gln Asp Ile Pro Ala Met Gln Gln Leu Ser Glu Trp Leu Met Lys Asn Leu Pro Asp Asn Asp Asn Glu Glu Asn Leu Ile His Gly Asp Phe Arg Leu Asp Asn 11e Val Phe His Pro Lys Glu Cys Arg Val Ile Ala Val Leu Asp Trp Glu Leu Ser Thr Ile

Gly His Pro Leu Ser Asp Leu Ala His Phe Ser Leu Phe Tyr Phe Trp Pro Arg Thr Val Pro Met 11e Asn Gln Gly Ser Tyr Ser Glu Asn Ser Gly lle Pro Ser Met Glu Glu Leu lle Ser Ile Tyr Cys Arg Cys Arg

Gly	Ile	Asn	Ser	He	Leu	Pro	Asn	Trp	Asn	Phe	Phe	Leu	Ala	Leu	Ser
			180					185					190		
Tyr	Phe	Lys	Met	Ala	Gly	Ile	Ala	Gln	Gly	Val	Tyr	Ser	Arg	Tyr	Leu
		195					200					205			
Leu	Gly	Asn	Asn	Ser	Ser	Glu	Asp	Ser	Phe	Leu	Phe	Ala	Asn	He	Val
	210					215					220				
Gln	Pro	Leu	Ala	Glu	Thr	Gly	Leu	Gln	Leu	Ser	Lys	Arg	Thr	Phe	Ser
225					230					235					240
Thr	Val	Leu	Pro	Gln	Ile	Asp	Thr	Thr	Gly	Gln	Leu	Phe	Val	Gln	Thr
				245					250					255	
Arg	Lys	Gly	Gln	Glu	Val	Leu	Ile	Lys	Val	Lys	His	Phe	Met	Lys	Gln
			260					265					270		
His	He	Leu	Pro	Ala	Glu	Lys	Glu	Val	Thr	Glu	Phe	Tyr	Val	G1n	Asn
		275					280					285			
Glu	Asn	Ser	Val	Asp	Lys	Trp	Gly	Lys	Pro	Leu	Val	11e	Asp	Lys	Leu
	290					295					300				
Lys	Glu	Met	Ala	Lys	Val	Glu	Gly	Leu	Trp	Asn	Leu	Phe	Leu	Pro	Ala
305					310					315					320
Val	Ser	Gly	Leu	Ser	His	Val	Asp	Tyr	Ala	Leu	Ile	Ala	Glu	Glu	Thṛ
				325					330					335	
Gly	Lys	Cys	Phe	Phe	Ala	Pro	Asp	Val	Phe	Asn	Cys	Gln	Ala	Pro	Лsp
			340					345					350		
Thr	Gly	Asn	Met	Glu	Val	Leu	His	Leu	Tyr	Gly	Ser	Glu	Glu	Gln	Lys
		355					360					365			
Lys	Gln	Trp	Leu	Glu	Pro	Leu	Leu	Gln	Gly	Asn	He	Thr	Ser	Cys	Phe
	370					375					380				
Cys	Met	Thr	Glu	Pro	Asp	Val	Ala	Ser	Ser	Asp	Ala	Thr	Asn	He	Glu
385					390					395					400
Cys	Ser	He	Gln	Arg	Asp	Glu	Asp	Ser	Tyr	Val	Пe	Asn	Gly	Lys	Lys
				405					410					415	
Trp	Trp	Ser	Ser	Gly	Ala	Gly	Asn	Pro	Lys	Cys	Lys	lle	Ala	He	Val
			420					425					430		
Leu	Gly	Arg	Thr	Gln	Asn	Thr	Ser	Leu	Ser	Arg					
		435					440								

```
<210> 4764
<211> 370
<212> PRT
<213> Homo sapiens
<400> 4764
Met Met Thr Pro Gln Val Ile Thr Pro Gln Gln Met Gln Gln Ile Leu
 1
                5
                                 10
Gln Gln Gln Val Leu Ser Pro Gln Gln Leu Gln Ala Leu Leu Gln Gln
           20
                             25
                                               30
Gln Gln Ala Val Met Leu Gln Gln Asp Phe Leu Asp Ser Gly Leu Glu
Asn Phe Arg Ala Ala Leu Glu Lys Asn Gln Gln Leu Gln Glu Phe Tyr
                      55
                                        60
Lys Lys Gln Gln Glu Gln Leu His Leu Gln Leu Leu Gln Gln Gln Gln
                   70
                                     75
65
90
               85
100
                             105
                                               110
Gln Gln Gln Gln His Pro Gly Lys Gln Ala Lys Glu Gln Gln Gln Gln
                         120
                                           125
Gln Gln Gln Gln Gln Leu Ala Ala Gln Gln Leu Val Phe Gln Gln
                     135
                                       140
Gln Leu Leu Gln Met Gln Gln Leu Gln Gln Gln His Leu Leu Ser
                  150
                                    155
                                                      160
145
Leu Gln Arg Gln Gly Leu Ile Ser 11e Pro Pro Gly Gln Ala Ala Leu
                                170
              165
Pro Val Gln Ser Leu Pro Gln Ala Gly Leu Ser Pro Ala Glu Ile Gln
                             185
                                               190
          180
Gln Leu Trp Lys Glu Val Thr Gly Val His Ser Met Glu Asp Asn Gly
                                           205
                         200
lle Lys His Gly Gly Leu Asp Leu Thr Thr Asn Asn Ser Ser Ser Thr
   210
                     215
                                       220
Thr Ser Ser Asn Thr Ser Lys Ala Ser Pro Pro Ile Thr His His Ser
```

Ile Val Asn Gly Gln Ser Ser Val Leu Ser Ala Arg Arg Asp Ser Ser 250 Ser His Glu Glu Thr Gly Ala Ser His Thr Leu Tyr Gly His Gly Val 260 265 270 Cys Lys Trp Pro Gly Cys Glu Ser Ile Cys Glu Asp Phe Gly Gln Phe 280 285 Leu Lys His Leu Asn Asn Glu His Ala Leu Asp Asp Arg Ser Thr Ala 295 300 Gln Cys Arg Val Gln Met Gln Val Val Gln Gln Leu Glu Ile Gln Leu 305 310 315 320 Ser Lys Glu Arg Glu Arg Leu Gln Ala Met Met Thr His Leu His Met 330 325 Arg Pro Ser Glu Pro Lys Pro Ser Pro Lys Pro Lys Leu Pro Gln Thr 350 345 Met Asn Phe Ile Lys Met Gln Met Ser Asp Leu His Leu Leu Met Gln 355 365 360 Leu Ser 370

<210> 4765

<211> 379

<212> PRT

<213> Homo sapiens

<400> 4765

Met Ser Arg Gly Tyr Ser Glu Asn Asn Asn Phe Leu Asn Asn Asn 1 5 10 15

Gln Met Val Leu Asp Met Ile Leu Tyr Pro Leu Ile Gly Ile Pro Gln 20 25 30

Thr lle Asn Trp Glu Thr lle Ala Arg Leu Val Pro Gly Leu Thr Pro
35 40 45

Lys Glu Cys Ala Lys Arg Phe Asp Glu Leu Lys Ser Ser Gly Ser Ser 50 55 60

Pro Val Asp Asn Gln Tyr Asn Ser Leu Met Ala Ala Gly Glu Ser Pro 65 70 75 80

Val	Glu	Thr	Leu	Ala	Thr	Tyr	lle	Lys	Ser	Ser	Leu	Leu	Asp	lle	His
				85					90					95	
Gly	Glu	Phe	Gln	Glu	Thr	Pro	Val	Gly	His	Asp	Ala	Val	Ser	Lys	Thr
			100					105					110		
Gly	Arg	His	Ser	11e	Ala	Ser	Thr	Arg	Λsn	Cys	Ser	Ser	Glu	Ser	Glu
		115					120					125			
Asn	Cys	Thr	Thr	His	Asn	Gly	Gly	Glu	Met	Thr	Glu	${\tt Glu}$	Ser	Glu	Gly
	130					135					140				
Pro	Asn	Met	Val	lle	His	Val	Cys	Asp	Glu	Ala	Lys	Asn	Leu	Lys	Glu
145					150					155					160
Лѕр	Phe	Thr	Cys	Pro	Arg	Asp	Leu	Leu	lle	Ser	Glu	Met	Lys	Tyr	Phe
				165					170					175	
Ala	Glu	Tyr	Leu	Ser	Met	Asp	Ala	Gln	Arg	Trp	Glu	Glu	Val	Asp	He
			180					185					190		
Ser	Val	His	Cys	Asp	Val	His	lle	Phe	Asn	Trp	Leu	lle	Lys	Tyr	lle
		195					200					205			
Lys	Arg	Asn	Thr	Lys	Glu	Asn	Lys	Asp	Cys	Glu	Met	Pro	Thr	Leu	Glu
	210					215					220				
Pro	Gly	Asn	Val	lle	Ser	Ile	Leu	Ile	Ser	Ser	Glu	Phe	Leu	Lys	Met
225					230					235					240
Asp	Ser	Leu	Val	Glu	Gln	Cys	lle	Gln	Tyr	Cys	His	Lys	Asn	Met	Asn
				245					250					255	
Ala	He	Val	Ala	Thr	Pro	Cys	Asn	Met	Asn	Cys	He	Asn	Ala	Asn	Leu
			260					265					270		
Leu	Thr	Arg	lle	Ala	Asp	Leu	Phe	Ser	His	Asn	Glu	Val	Asp	Asp	Leu
		275					280					285			
Lys	Asp	Lys	Lys	Asp	Lys	Phe	Lys	Ser	Lys	Leu	Phe	Cys	Lys	Lys	He
	290					295					300				
Glu	Arg	Leu	Phe	Asp	Pro	Glu	Tyr	Leu	Asn	Pro	Asp	Ser	Arg	Ser	
305					310					315					320
Ala	Ala	Thr	Leu	Tyr	Arg	Cys	Cys	Leu	Cys	Lys	Lys	Leu	Leu		Lys
				325					330					335	
Glu	Thr	G] u	Arg	Arg	lle	Pro	Cys		Pro	G1 y	Lys	lle		Val	Asp
			340					345					350		
Arg	Arg		Asn	He	Val	Tyr		His	Пе	Arg	Cys		Glu	Asp	Lys
		355					360					365			

11e His Thr Cys Ile Phe Val Tyr 11e Tyr 11e 370 375

<210> 4766

<211> 155

<212> PRT

<213> Homo sapiens

<400> 4766

Met His Lys Tyr Ile Leu Ile Val Lys Asp Leu Asn Pro Thr Glu Pro

1 5 10 15

Pro Gln Cys Val Gly 11e Phe Pro Pro Leu Ser Pro 11e His Ala Ser 20 25 30

Leu Leu Tyr Pro Leu Pro Pro Cys Ser Gly Pro Pro Cys Pro Leu Ser 35 40 45

Ser Cys Trp Val Cys Cys Trp Glu Ser Ala Gly Asp Arg Lys Asp Arg
50 55 60

Gly Glu Arg Gly Gln Gly Thr Asn Phe Pro Ser Pro Pro Leu Gln Ser
65 70 75 80

His Gly Pro Ala Ala Phe Leu Tyr Gln Arg Phe Gln Leu Leu Pro Gly
85 90 95

Gly Val Pro Lys Ser Ser Phe Ser Leu Gly His His Ala Ser Leu Pro 100 105 110

Pro Ser Gln Pro Lys Ala Ser His Gly Ala His Arg Phe Leu Ser Arg 115 120 125

Val Leu Ser 11e Met Phe Leu Cys 11e Leu Ser Met Leu Tyr Asn Trp 130 135 140

Ser Phe 11e Lys Phe Cys Leu Asn Gly Pro 11e 145 150 155

<210> 4767

<211> 421

<212> PRT

<213> Homo sapiens

<400)> 47	767													
Met	Met	Tyr	Arg	Thr	Val	Gly	Phe	Gly	Thr	Arg	Ser	Arg	Asn	Leu	Lys
1				5					10					15	
Pro	Trp	Met	He	Ala	Val	Leu	Пe	Val	Leu	Ser	Leu	Thr	Val	Val	Ala
			20					25					30		
Val	Thr	He	Gly	Leu	Leu	Val	His	Phe	Leu	Val	Phe	Asp	Gln	Lys	Lys
		35					40					45			
Glu	Tyr	Tyr	His	Gly	Ser	Phe	Lys	lle	Leu	Asp	Pro	Gln	lle	Asn	Asn
	50					55					60				
Asn	Phe	Gly	Gln	Ser	Asn	Thr	Tyr	Gln	Leu	Lys	Asp	Leu	Arg	Glu	Thr
65					70					75					80
Thr	Glu	Asn	Leu	Val	Ser	Gln	Val	Asp	Glu	lle	Phe	He	Asp	Ser	Ala
				85					90					95	
Trp	Lys	Lys	Asn	Tyr	lle	Lys	Asn	Gln	Val	Val	Arg	Leu	Thr	Pro	Glu
			100					105					110		
Glu	Asp		Val	Lys	Val	Asp		lle	Met	Val	Phe		Phe	Pro	Ser
		115					120					125			
Thr		Gln	Arg	Ala	Val	Arg	Glu	Lys	Lys	He		Ser	lle	Leu	Asn
	130					135					140		_		
	Lys	He	Arg	Asn		Arg	Ala	Leu	Pro		Asn	Ala	Ser	Ser	
145					150			mı.	0.1	155	,	mı		0.1	160
GIn	Val	Asn	Ala		Ser	Ser	Ser	Thr		Glu	Leu	lhr	Val		Ala
	0	0.1		165		17 1	13	,	170					175	C
Ser	Cys	61y		Arg	Val	Val	Pro		Asn	Val	Asn	Arg		Ala	Ser
C1	17 J	7.1	180	n	1	A 7	41.	185	D	Т	C1	41 -	190	1	C1
61 y	vai		Ala	Pro	Lys	Ala		ırp	rro	тр	GIN		ser	Leu	GIN
т	Aan	195	11.	Ша	Cla	Cva	200	A1 o	The	Lou	110	205	Acn	Thr	Trn
Tyr	210	ASII	1.16	шѕ	GIII	Cys 215	Giy	MIA	1111	Leu	220	361	VSII	1111	пþ
Lon		Thr	Λla	Ala	Hic	Cys	Pho	Cln	Lve	Tyr		Acn	Pro	Hic	Gln
225	vai	1111	Ма	Ма	230	Cys	1116	0111	Lys	235	rivs	ASII	110	1115	240
	The	Val	Sor	Phe		Thr	Lve	He	Asn		Pro	Lau	Met	Lve	
пр	1 117	, 41	061	245	Ory	1111	ra y co	110	250			i, cu	inc t	255	5
Asn	Val	Arø	Arø		He	lle	Hic	Glu		Tvr	Arø	Ser	Λla		Arø
11011		6	260	7 710	. 10	.10	1113	265	ى ر ـ ـ	- , 1	5	501	270	0	8

Glu Tyr Asp lle Ala Val Val Gln Val Ser Ser Arg Val Thr Phe Ser 280 Asp Asp Ile Arg Arg Ile Cys Leu Pro Glu Ala Ser Ala Ser Phe Gln 290 295 300 Pro Asn Leu Thr Val His Ile Thr Gly Phe Gly Ala Leu Tyr Tyr Gly 315 305 310 Gly Glu Ser Gln Asn Asp Leu Arg Glu Ala Arg Val Lys Ile Ile Ser 325 330 Asp Asp Val Cys Lys Gln Pro Gln Val Tyr Gly Asn Asp Ile Lys Pro 350 340 345 Gly Met Phe Cys Ala Gly Tyr Met Glu Gly Ile Tyr Asp Ala Cys Arg 360 365 Gly Asp Ser Gly Gly Pro Leu Val Thr Arg Asp Leu Lys Asp Thr Trp 375 380 Tyr Leu Ile Gly Ile Val Ser Trp Gly Asp Asn Cys Gly Gln Lys Asp 385 390 395 400 Lys Pro Gly Val Tyr Thr Gln Val Thr Tyr Tyr Arg Asn Trp lle Ala 410 415 Ser Lys Thr Gly Ile 420

<210> 4768

<211> 495

<212> PRT

<213> Homo sapiens

<400> 4768

Met Ala Ala Glu Pro Gln Pro Ser Ser Leu Ser Tyr Arg Thr Thr Gly

1 5 10 15

Ser Thr Tyr Leu His Pro Leu Ser Glu Leu Leu Gly Ile Pro Leu Asp
20 25 30

Gln Val Asn Phe Val Val Cys Gln Leu Val Ala Leu Phe Ala Ala Phe
35 40 45

Trp Phe Arg 11e Tyr Leu Arg Pro Gly Thr Thr Ser Ser Asp Val Arg
50 55 60

His	Ala	Val	Ala	Thr	He	Phe	Gly	He	Tyr	Phe	Val	lle	Phe	Cys	Phe
65					70					75					80
Gly	Trp	Tyr	Ser	Val	His	Leu	Phe	Val	Leu	Val	Leu	Met	Cys	Tyr	Ala
				85					90					95	
He	Met	Val	Thr	Ala	Ser	Val	Ser	Asn	He	His	Arg	Tyr	Ser	Phe	Phe
			100					105					110		
Val	Ala	Met	Gly	Tyr	Leu	Thr	He	Cys	His	He	Ser	Arg	He	Tyr	He
		115					120					125			
Phe	His	Tyr	Gly	Ile	Leu	Thr	Thr	Asp	Phe	Ser	Gly	Pro	Leu	Met	He
	130					135					140				
Val	Thr	Gln	Lys	Ile	Thr	Thr	Leu	Ala	Phe	Gln	Val	His	Asp	Gly	Leu
145					150					155					160
Gly	Arg	Arg	Ala	Glu	Asp	Leu	Ser	Ala	Glu	Gln	His	Arg	Leu	Ala	He
				165					170					175	
Lys	Val	Lys	Pro	Ser	Phe	Leu	Glu	Tyr	Leu	Ser	Tyr	Leu	Leu	Asn	Phe
			180					185					190		
Met	Ser	Val	Ile	Ala	Gly	Pro	Cys	Asn	Asn	Phe	Lys	Asp	Tyr	He	Ala
		195					200					205			
Phe	Ile	Glu	Gly	Lys	His	Ile	His	Met	Lys	Leu	Leu	Glu	Val	Asn	Trp
	210					215					220				
Lys	Arg	Lys	Gly	Phe	His	Ser	Leu	Pro	Glu	Pro	Ser	Pro	Thr	Gly	Ala
225					230					235					240
Val	lle	His	Lys	Leu	Gly	He	Thr	Leu	Val	Ser	Leu	Leu	Leu	Phe	Leu
				245					250					255	
Thr	Leu	Thr	Lys	Thr	Phe	Pro	Val	Thr	Cys	Leu	Val	Asp	Asp	Trp	Phe
			260					265					270		
Val	His	Lys	Ala	Ser	Phe	Pro	Ala	Arg	Leu	Cys	Tyr	Leu	Tyr	Val	Val
		275					280					285			
Met	Gln	Ala	Ser	Lys	Pro	Lys	Tyr	Tyr	Phe	Ala	Trp	Thr	Leu	Ala	Asp
	290					295					300				
Ala	Val	Asn	Asn	Ala	Ala	Gly	Phe	Gly	Phe	Ser	Gly	Val	Asp	Lys	Asn
305					310					315					320
Gly	Asn	Phe	Cys	Trp	Asp	Leu	Leu	Ser	Asn	Leu	Asn	11e	Trp	Lys	lle
				325					330					335	
Glu	Thr	Ala	Thr	Ser	Phe	Lys	Met	Tyr	Leu	Glu	Asn	Trp	Asn	He	Gln
			340					345					350		

Thr Ala Thr Trp Leu Lys Cys Val Cys Tyr Gln Arg Val Pro Trp Tyr Pro Thr Val Leu Thr Phe Ile Leu Ser Ala Leu Trp His Gly Val Tyr Pro Gly Tyr Tyr Phe Thr Phe Leu Thr Gly Ile Leu Val Thr Leu Ala Ala Arg Ala Val Arg Asn Asn Tyr Arg His Tyr Phe Leu Ser Ser Arg Ala Leu Lys Ala Val Tyr Asp Ala Gly Thr Trp Ala Val Thr Gln Leu Ala Val Ser Tyr Thr Val Ala Pro Phe Val Met Leu Ala Val Glu Pro Thr Ile Ser Leu Tyr Lys Ser Met Tyr Phe Tyr Leu His Ile Ile Ser Leu Leu Ile Ile Leu Phe Leu Pro Met Lys Pro Gln Ala His Thr Gln Arg Arg Pro Gln Thr Leu Asn Ser lle Asn Lys Arg Lys Thr Asp

<210> 4769

<211> 953

<212> PRT

<213> Homo sapiens

<400> 4769

Met Cys Asn Pro Glu Glu Ala Ala Leu Leu Arg Leu Glu Glu Val Phe 1 $$ $$

65					70					75					80
Thr	Gly	Leu	Glu	Ser	Leu	Leu	Leu	Leu	Arg	Gly	Ala	Λsp	Arg	Val	Leu
				85					90					95	
Gln	Ala	His	lle	Glu	Tyr	11e	Glu	Ser	Tyr	Thr	Ser	Cys	Met	Val	Val
			100					105					110		
Gln	Ala	Phe	Gln	Lys	Ala	Ala	Lys	Arg	Arg	Ser	Glu	Tyr	Trp	Arg	Gly
		115					120					125			
Gln	Arg	Lys	Ala	Leu	Arg	Gln	Leu	Leu	Ser	Gly	Val	Ser	Ser	Glu	Gly
	130					135					140				
Ser	Val	Gly	Ala	Ser	Leu	Gly	Gln	Ala	Leu	His	Gln	Pro	Leu	Ala	His
145					150					155					160
His	Val	Gln	Gln	Tyr	Val	Leu	Leu	Leu	Leu	Ser	Pro	Gly	Asp	Thr	lle
				165					170					175	
G1 y	Glu	His	His	Pro	Thr	Arg	Glu	Leu	Val	Val	Asn	Ala	Val	Thr	Leu
			180					185					190		
Phe	Gly	Asn	Leu	Gln	Ser	Phe	Met	Lys	Gln	Glu	Leu	Asp	Gln	Ala	Val
		195					200					205			
Ala	Thr	Gln	Ala	Leu	Trp	His	Thr	Leu	Arg	Gly	Arg	Leu	Arg	Asp	Val
	210					215					220				
Leu	Cys	Thr	Pro	Ala	His	Arg	Leu	Leu	Gln	Asp	Ser	Gln	Asp	Val	Pro
225					230					235					240
Val	Thr	Val	Ala	Pro	Leu	Arg	Ala	Glu	Arg	Val	Leu	Leu	Phe	Asp	Asp
				245					250					255	
Ala	Leu	Val	Leu	Leu	Gln	Gly	His	Asn	Val	His	Thr	Phe	Asp	Leu	Lys
			260					265					270		
Leu	Val	Trp	Val	Asp	Pro	G1 y	G1n	Asp	Gly	Cys	Thr	Phe	His	Leu	Leu
		275										285			
Thr		Glu	Glu	Glu	Phe		Phe	Cys	Ala	Lys		Ser	Gln	G1 y	Gln
	290					295		_			300				
	Val	Trp	Gln	Trp		Val	Thr	Trp	Ala		His	GIn	Ala	Leu	
305					310					315	_ ^				320
Gly	Lys	Lys	Asp		Pro	Val	Leu	Gly		G1 y	Leu	Glu	Pro		GIn
				325					330		- "			335	
Pro	Pro	Asp		Arg	Cys	Ala	Glu		Thr	Phe	GIn	Ala		G1 y	Arg
	0	0.3	340	Tr.	T	61	61	345	Tr.	0		<i>C</i> ³	350	15	
Leu	Cys	GIn	Ala	Ihr	Lyr	Glu	Gly	Glu	Trp	Cys	Arg	Ыγ	Arg	Pro	HIS

		355					360					365			
Gly	Lys	Gly	Thr	Leu	Lys	Trp	Pro	Asp	Gly	Arg	Asn	His	Val	Gly	Asn
	370					375					380				
Phe	Cys	Gln	Gly	Leu	Glu	His	Gly	Phe	Gly	Пe	Arg	Leu	Leu	Pro	Gln
385					390					395					400
Ala	Ser	Glu	Asp	Lys	Phe	Asp	Cys	Tyr	Lys	Cys	His	Trp	Arg	Glu	Gly
				405					410					415	
Ser	Met	Cys	Gly	Tyr	Gly	Пe	Cys	Glu	Tyr	Ser	Thr	Asp	Gly	Val	Tyr
			420					425					430		
Lys	Gly	Tyr	Phe	Gln	Glu	Gly	Leu	Arg	His	Gly	Phe	Gly	Val	Leu	Glu
		435					440					445			
Ser	Gly	Pro	Gln	Ala	Pro	Gln	Pro	Phe	Arg	Tyr	Thr	Gly	His	Trp	Glu
	450					455					460				
Arg	Gly	Gln	Arg	Ser	Gly	Tyr	Gly	He	Glu	Glu	Asp	Gly	Asp	Arg	Gly
465					470					475					480
Glu	Arg	Tyr	lle	Gly	Met	Trp	Gln	Ala	Gly	Gln	Arg	His	Gly	Pro	Gly
				485					490					495	
Val	Met	Val	Thr	Gln	Ala	Gly	Val	Cys	Tyr	Gln	Gly	Thr	Phe	Gln	Ala
			500					505					510		
		Thr	500		Pro	Gly	lle		Leu	Ser	G1u	Asp		Ser	Leu
		Thr 515	500		Pro	Gly	11e 520		Leu	Ser	G1u	Asp 525		Ser	Leu
Asp	Lys	515	500 Val	Gly	Pro Thr		520	Leu				525	Asp		
Asp	Lys	515	500 Val	Gly			520	Leu				525	Asp		
Asp Tyr	Lys Glu 530	515 Gly	500 Val Thr	G1y Phe		Arg 535	520 Asp	Leu Leu	Thr	Leu	Met 540	525 Gly	Asp Lys	Gly	Lys
Asp Tyr	Lys Glu 530	515 Gly	500 Val Thr	G1y Phe	Thr	Arg 535	520 Asp	Leu Leu	Thr	Leu	Met 540	525 Gly	Asp Lys	Gly	Lys
Asp Tyr Val 545	Lys Glu 530 Thr	515 Gly Phe	500 Val Thr Pro	Gly Phe Asn	Thr Gly	Arg 535 Phe	520 Asp Thr	Leu Leu Leu	Thr Glu	Leu Gly 555	Met 540 Ser	525 G1y Phe	Asp Lys Gly	Gly Ser	Lys Gly 560
Asp Tyr Val 545	Lys Glu 530 Thr	515 Gly Phe	500 Val Thr Pro	Gly Phe Asn	Thr G1y 550	Arg 535 Phe	520 Asp Thr	Leu Leu Leu	Thr Glu Val	Leu Gly 555	Met 540 Ser	525 G1y Phe	Asp Lys Gly	Gly Ser	Lys Gly 560 Leu
Asp Tyr Val 545 Ala	Lys Glu 530 Thr	515 Gly Phe Arg	500 Val Thr Pro Gly	Gly Phe Asn Leu 565	Thr Gly 550 His	Arg 535 Phe Thr	520 Asp Thr Gln	Leu Leu Leu Gly	Thr Glu Val 570	Leu Gly 555 Leu	Met 540 Ser Asp	525 Gly Phe Thr	Asp Lys Gly Ala	Gly Ser Ala 575	Lys Gly 560 Leu
Asp Tyr Val 545 Ala Pro	Lys Glu 530 Thr Gly Pro	515 Gly Phe Arg Asp	500 Val Thr Pro Gly Pro 580	Gly Phe Asn Leu 565 Ser	Thr Gly 550 His	Arg 535 Phe Thr	520 Asp Thr Gln Cys	Leu Leu Gly Lys 585	Thr Glu Val 570 Arg	Leu Gly 555 Leu Gln	Met 540 Ser Asp	525 Gly Phe Thr	Asp Lys Gly Ala Val 590	Gly Ser Ala 575 Gly	Lys Gly 560 Leu Ala
Asp Tyr Val 545 Ala Pro	Lys Glu 530 Thr Gly Pro	515 Gly Phe Arg Asp	500 Val Thr Pro Gly Pro 580	Gly Phe Asn Leu 565 Ser	Thr Gly 550 His	Arg 535 Phe Thr	520 Asp Thr Gln Cys	Leu Leu Gly Lys 585	Thr Glu Val 570 Arg	Leu Gly 555 Leu Gln	Met 540 Ser Asp	525 Gly Phe Thr Gly	Asp Lys Gly Ala Val 590	Gly Ser Ala 575 Gly	Lys Gly 560 Leu
Asp Tyr Val 545 Ala Pro	Lys Glu 530 Thr Gly Pro	515 Gly Phe Arg Asp Val 595	500 Val Thr Pro Gly Pro 580 Glu	Gly Phe Asn Leu 565 Ser	Thr Gly 550 His Ser	Arg 535 Phe Thr Thr	520 Asp Thr Gln Cys Gln 600	Leu Leu Gly Lys 585 Gly	Thr Glu Val 570 Arg Val	Leu Gly 555 Leu Gln Tyr	Met 540 Ser Asp Leu Ser	525 Gly Phe Thr Gly Pro 605	Asp Lys Gly Ala Val 590 Phe	Gly Ser Ala 575 Gly Arg	Lys Gly 560 Leu Ala
Asp Tyr Val 545 Ala Pro	Lys Glu 530 Thr Gly Pro Pro	515 Gly Phe Arg Asp Val 595	500 Val Thr Pro Gly Pro 580 Glu	Gly Phe Asn Leu 565 Ser	Thr Gly 550 His	Arg 535 Phe Thr Thr	520 Asp Thr Gln Cys Gln 600	Leu Leu Gly Lys 585 Gly	Thr Glu Val 570 Arg Val	Leu Gly 555 Leu Gln Tyr	Met 540 Ser Asp Leu Ser Glu	525 Gly Phe Thr Gly Pro 605	Asp Lys Gly Ala Val 590 Phe	Gly Ser Ala 575 Gly Arg	Lys Gly 560 Leu Ala
Asp Tyr Val 545 Ala Pro Phe	Lys Glu 530 Thr Gly Pro Pro Val 610	515 Gly Phe Arg Asp Val 595 Cys	500 Val Thr Pro Gly Pro 580 Glu	Gly Phe Asn Leu 565 Ser Ser	Thr Gly 550 His Ser Arg Cys	Arg 535 Phe Thr Thr Pro 615	520 Asp Thr Gln Cys Gln 600 Arg	Leu Leu Gly Lys 585 Gly Asp	Thr Glu Val 570 Arg Val	Leu Gly 555 Leu Gln Tyr	Met 540 Ser Asp Leu Ser Glu 620	525 Gly Phe Thr Gly Pro 605 Ala	Asp Lys Gly Ala Val 590 Phe Leu	Gly Ser Ala 575 Gly Arg Leu	Lys Gly 560 Leu Ala Asp Gly
Asp Tyr Val 545 Ala Pro Phe Phe	Lys Glu 530 Thr Gly Pro Pro Val 610	515 Gly Phe Arg Asp Val 595 Cys	500 Val Thr Pro Gly Pro 580 Glu	Gly Phe Asn Leu 565 Ser Ser	Thr Gly 550 His Ser Arg Cys	Arg 535 Phe Thr Thr Pro 615	520 Asp Thr Gln Cys Gln 600 Arg	Leu Leu Gly Lys 585 Gly Asp	Thr Glu Val 570 Arg Val	Leu Gly 555 Leu Gln Tyr Gln Arg	Met 540 Ser Asp Leu Ser Glu 620	525 Gly Phe Thr Gly Pro 605 Ala	Asp Lys Gly Ala Val 590 Phe Leu	Gly Ser Ala 575 Gly Arg Leu	Lys Gly 560 Leu Ala Asp Gly Leu
Asp Tyr Val 545 Ala Pro Phe Phe 625	Lys Glu 530 Thr Gly Pro Val 610 Asp	515 Gly Phe Arg Asp Val 595 Cys	500 Val Thr Pro Gly Pro 580 Glu Ala Gln	Gly Phe Asn Leu 565 Ser Gly Ser	Thr Gly 550 His Ser Arg Cys	Arg 535 Phe Thr Thr Pro 615 Arg	520 Asp Thr Gln Cys Gln 600 Arg	Leu Leu Gly Lys 585 Gly Asp Leu	Thr Glu Val 570 Arg Val Leu Arg	Leu Gly 555 Leu Gln Tyr Gln Arg 635	Met 540 Ser Asp Leu Ser Glu 620 Ser	525 Gly Phe Thr Gly Pro 605 Ala	Asp Lys Gly Ala Val 590 Phe Leu	Gly Ser Ala 575 Gly Arg Leu Tyr	Lys Gly 560 Leu Ala Asp Gly Leu 640

				645					650					655	
Ile	Leu	Glu	Glu	Leu	Leu	Gln	His	Arg	Glu	Pro	Lys	Ala	Leu	Gln	Leu
			660					665					670		
Tyr	Leu	Arg	Lys	Ala	Leu	Ser	Asn	Ser	Leu	His	Pro	Leu	G1 y	Lys	Leu
		675					680					685			
Leu	Arg	Thr	Leu	Met	Leu	Thr	Phe	Gln	Ala	Thr	Tyr	Ala	Gly	Val	Gly
	690					695					700				
Ala	Asn	Lys	His	Leu	Gln	Glu	Leu	Ala	Gln	Glu	Glu	Val	Lys	Gln	His
705					710					715					720
Ala	Gln	Glu	Leu	Trp	Ala	Ala	Tyr	Arg	Gly	Leu	Leu	Arg	Val	Ala	Leu
				725					730					735	
Glu	Arg	Lys	Gly	Gln	Ala	Leu	Glu	Glu	Asp	Glu	Asp	Thr	Glu	Thr	Arg
			740					745					750		
Asp	Leu	Gln	Val	His	Gly	Leu	Val	Leu	Pro	Leu	Met	Leu	Pro	Ser	Phe
		755					760					765			
Tyr	Ser	Glu	Leu	Phe	Thr	Leu	Tyr	Leu	Leu	Leu	His	Glu	Arg	Glu	Asp
	770					775					780				
Ser	Phe	Tyr	Ser	Gln	Gly	Ile	Ala	Asn	Leu	Ser	Leu	Phe	Pro	Asp	Thr
785					790					795					800
Gln	Leu	Leu	Glu	Phe	Leu	Asp	Val	Gln	Lys	His	Leu	Trp	Pro	Leu	Lys
				805					810					815	
Asp	Leu	Thr	Leu	Thr	Ser	Asn	Gln	Arg	Tyr	Ser	Leu	Val	Arg	Asp	Lys
			820					825					830		
Cys	Phe	Leu	Ser	Ala	Thr	Glu	Cys	Leu	Gln	Lys	lle	Met	Thr	Thr	Val
		835					840					845			
Asp		Arg	Glu	Lys	Leu	Glu	Val	Leu	Glu	Arg		Tyr	Gly	Glu	He
	850					855					860				
	Gly	Thr	Val	Ser		Val	l.eu	Gly	Arg		Tyr	Lys	Leu	Pro	
865			_	_	870					875	_				880
Asp	Λsp	Leu	Leu		Leu	Leu	He	Tyr		Val	Ser	His	Ala		He
			0.1	885	0.1			•	890					895	D
GIn	His	Leu		Ala	Glu	He	H1S		He	Arg	Asp	Met		Asp	Pro
		TC I	900	0.1		T		905	,		Ti	. 1	910	C1	C
Asn	ніѕ		ыу	ыу	Leu	Tyr		rne	Leu	Leu	ınr		Leu	61 u	ser
	-	915	n	71.	C1.	,	920			Α		925	Α		n
Cvs	177	4 . 1					1.111	Acr	Mot			H17 ^		יים	Pro

930 935 940 Gly His Trp His Ser Arg Glu Leu Trp 945 950

<210> 4770

<211> 331

<212> PRT

<213> Homo sapiens

<400> 4770

Met Asn Val Lys Asn Val Gly Lys Leu Asn Gln Ser Ser Asp Leu Leu

1 5 10 15

Arg His His Arg 11e His Ser Gly Glu Lys Pro Tyr Val Cys Asn Lys
20 25 30

Cys Gly Glu Ser Phe Arg Ser Ser Ser Asp Leu Ile Lys His His Arg
35 40 45

Val His Thr Gly Glu Lys Pro His Glu Cys Ser Glu Cys Gly Lys Val
50 55 60

Phe Ser Gln Arg Ser His Leu Val Thr His Gln Lys 11e His Thr Gly
65 70 75 80

Glu Lys Pro Tyr Gln Cys Thr Glu Cys Glu Lys Ala Phe Arg Arg Arg

85 90 95

Ser Leu Leu lle Gln Arg Arg 11e His Ser Gly Glu Lys Pro Cys 100 105 110

Glu Cys Lys Glu Cys Gly Lys Leu Phe Met Trp His Thr Ala Phe Leu

Lys His Gln Arg Leu His Ala Gly Glu Lys Leu Glu Glu Cys Glu Lys 130 135 140

Thr Phe Ser Lys Asp Glu Glu Leu Arg Gly Glu Gln Lys Ile His Gln 145 150 155 160

Glu Glu Lys Ala Tyr Trp Cys Asn Gln Cys Gly Arg Ala Phe Gln Gly

165 170 175

Ser Ser Asp Leu 11e Gly His Gln Val Thr His Thr Gly Glu Lys Pro 180 185 190

Tyr Glu Cys Lys Glu Cys Gly Lys Thr Phe Asn Gln Ser Ser Asp Leu

		195					200					205			
Leu	Arg	His	His	Arg	lle	His	Ser	Gly	Glu	Lys	Pro	Tyr	Val	Cys	Asn
	210					215					220				
Lys	Cys	Gly	Lys	Ser	Phe	Arg	Gly	Ser	Ser	Asp	Leu	11e	Arg	His	His
225					230					235					240
Arg	Val	His	Thr	Gly	Glu	Lys	Pro	Tyr	Glu	Cys	Pro	Glu	Cys	Trp	Lys
				245					250					255	
Ala	Phe	Ser	Gln	Asn	Ser	His	Leu	Val	Ser	His	Gln	Arg	lle	His	Thr
			260					265					270		
Arg	Glu	Lys	Pro	Phe	Glu	Cys	Ser	Asn	Cys	Gly	Lys	Ala	Phe	Ser	Gly
		275					280					285			
Trp	Thr	Ala	Phe	Leu	Lys	His	Gln	Lys	Leu	His	lle	Gly	Lys	Glu	Phe
	290					295					300				
Glu	Asp	Cys	Lys	Ser	Leu	Gln	Thr	G1 y	Pro	Пе	Leu	He	Gly	Ser	Arg
305					310					315					320
Asn	Leu	Met	Asn	Ala	Val	Lys	Leu	G1 y	Lys	Val					
				325					330						

<210> 4771

<211> 1099

<212> PRT

<213> Homo sapiens

<400> 4771

Met Ala His Leu Gly Pro Thr Pro Pro Pro His Ser Leu Asn Tyr Lys Ser Glu Asp Arg Leu Ser Glu Gln Asp Trp Pro Ala Tyr Phe Lys Val 20 25 30 Pro Cys Cys Gly Val Asp Thr Ser Gln Ile Glu Ser Glu Glu Ala Glu 35 40 45 Val Asp Val Arg Glu Arg Glu Thr Gln Arg Asp Arg Glu Pro Lys Arg 55 60 Ala Arg Asp Leu Thr Leu Arg Asp Ser Cys Thr Asp Asn Ser Met Gln 65 70 75 80 Phe Gly Thr Arg Thr Thr Ala Glu Pro Gly Phe Met Gly Thr Trp

					85					90					95	
G	ln	Asn	Ala	Asp	Thr	Asn	Leu	Leu	Phe	Arg	Met	Ser	Gln	Gln	Ala	Ile
				100					105					110		
٨ı	rg	Cys	Thr	Leu	Val	Asn	Cys	Thr	Cys	Glu	Cys	Phe	Gln	Pro	Gly	Lys
			115					120					125			
11	le	Asn	Leu	Arg	Thr	Cys	Asp	Gln	Cys	Lys	His	Gly	Trp	Val	Ala	His
		130					135					140				
A.	la	Leu	Asp	Lys	Leu	Ser	Thr	Gln	His	Leu	Tyr	His	Pro	Thr	Gln	Val
14	45					150					155	٠				160
G.	lu	He	Val	G1n	Ser	Asn	Val	Val	Phe	Asp	Ile	Ser	Ser	Leu	Met	Leu
					165					170					175	
T	yr	G1 y	Thr	Gln	Ala	Val	Pro	Val	Arg	Leu	Lys	He	Leu	Leu	Asp	Arg
				180					185					190		
Le	eu	Phe	Ser	Val	Leu	Lys	Gln	Glu	Glu	Val	Leu	His	He	Leu	His	Gly
			195					200					205			
Le	eu	Gly	Trp	Thr	Leu	Arg	Asp	Tyr	Val	Arg	Gly	Tyr	lle	Leu	Gln	Asp
		210					215					220				
A.	la	Ala	Gly	Lys	Val	Leu	Asp	Arg	Trp	Ala	Ile	Met	Ser	Arg	Glu	Glu
22	25					230					235					240
G.	lu	He	He	Thr	Leu	Gln	Gln	Phe	Leu	Arg	Phe	Gly	Glu	Thr	Lys	Ser
					245					250					255	
1	le	Val	Glu	Leu	Met	Ala	He	Gln	Glu	Lys	Glu	Gly	Gln	Ala	Val	Ala
				260					265					270		
Va	al	Pro	Ser	Ser	Lys	Thr	Asp	Ser	Asp	lle	Arg	Thr	Phe	He	Glu	Ser
			275					280					285			
As	sn	Asn	Arg	Thr	Arg	Ser	Pro	Ser	Leu	Leu	Ala	His	Leu	Glu	Asn	Ser
		290					295					300				
As	sn	Pro	Ser	Ser	lle	His	His	Phe	Glu	Asn	lle	Pro	Asn	Ser	Leu	
)5					310					315					320
Pł	1e	Leu	Leu	Pro		Gln	Tyr	He	Asn		Val	Ser	Ala	Pro	Leu	Leu
					325					330					335	
G.	lу	Leu	Pro		Asn	Gly	Leu	Leu		Glu	G1n	Pro	Gly		Arg	Leu
				340					345					350		
Aı	rg	Glu		Ser	Leu	Ser	Thr		Asn	Glu	Tyr	Asn		Ser	Ser	Glu
			355					360					365			

Ser	Glu 370	Val	Ser	Pro	Thr	Pro 375	Tyr	Lys	Asn	Asp	G1n 380	Thr	Pro	Asn	Arg
Asn	Ala	Leu	Thr	Ser	He	Thr	Λsn	Val	Glu	Pro	Lys	Thr	Glu	Pro	Ala
385					390					395					400
Cys	Val	Ser	Pro	He	Gln	Asn	Ser	Λla	Pro	Val	Ser	Asp	Leu	Thr	Lys
				405					410					415	
Thr	Glu	His	Pro	Lys	Ser	Ser	Phe	Arg	Ile	His	Arg	Met	Arg	۸rg	Met
			420					425					430		
Gly	Ser	Ala	Ser	Arg	Lys	Gly	Arg	Val	Phe	Cys	Asn	Ala	Cys	Gly	Lys
		435					440					445			
Thr	Phe	Tyr	Asp	Lys	Gly	Thr	Leu	Lys	Ile	His	Tyr	Asn	Ala	Val	His
	450					455					460				
Leu	Lys	Πe	Lys	His	Arg	Cys	Thr	Ile	Glu	Gly	Cys	Asn	Met	Val	Phe
465					470					475					480
Ser	Ser	Leu	Arg	Ser	Arg	Asn	Arg	His	Ser	Ala	Asn	Pro	Asn	Pro	Arg
				485					490					495	
Leu	His	Met	Pro	Met	Leu	Arg	Asn	Asn	Arg	Asp	Lys	Asp	Leu	lle	Arg
			500					505					510		
Ala	Thr	Ser	Gly	Ala	Ala	Thr	Pro	Val	lle	Ala	Ser	Thr	Lys	Ser	Asn
		515					520					525			
Leu	Ala	Leu	Thr	Ser	Pro	Gly	Arg	Pro	Pro	Met	Gly	Phe	Thr	Thr	Pro
	530					535					540				
Pro	Leu	Asp	Pro	Val	Leu	Gln	Asn	Pro	Leu	Pro	Ser	Gln	Leu	Val	Phe
545					550					555					560
Ser	Gly	Leu	Lys	Thr	Val	Gln	Pro	Val	Pro	Pro	Phe	Tyr	Arg	Ser	Leu
				565					570					575	
Leu	Thr	Pro	Gly	Glu	Met	Val	Ser	Pro	Pro	Thr	Ser	Leu	Pro	Thr	Ser
			580					585					590		
Pro	lle	He	Pro	Thr	Ser	Gly		He	Glu	Gln	His		Pro	Pro	Pro
		595					600					605			
Ser		Pro	Va]	Val	Pro		Val	Met	Met	Ala		His	Glu	Pro	Ser
	610					615	_				620				
	Asp	Leu	Ala	Pro		Lys	Lys	Pro	Arg		Ser	Ser	Met	Pro	
625					630			<i>m</i> :		635	0.3	13.1			640
Lys	He	Glu	Lys		He	He	Asp	Thr			Glu	Phe	Asp	Asp	Glu
				645					650					655	

Asp	Asp	Asp	Pro	Asn	Asp	Gly	Gly		Val	Val	Asn	Asp	Met	Ser	His
			660					665					670		
Asp	Asn	His 675	Cys	His	Ser	Gln	G1u 680	Glu	Met	Ser	Pro	Gly 685	Met	Ser	Val
Lys	Asp 690	Phe	Ser	Lys	His	Asn 695	Arg	Thr	Arg	Cys	11e 700	Ser	Arg	Thr	Glu
Tle		Aro	Ala	Asn	Ser		Thr	Ser	Glu	Asn		Glu	Pro	G1n	Aro
705	ni g	111 6	діа	пор	710	me c	1111	SCI	ora	715	OIII	oru	110	oru	720
Asp	Tyr	Glu	Asn	Glu	Ser	Glu	Ser	Ser	Glu	Pro	Lys	Leu	Gly	Glu	Glu
				725					730					735	
Ser	Met	Glu	Gly	Asp	Glu	His	Ile	His	Ser	Glu	Val	Ser	Glu	Lys	Val
			740					745					750		
Leu	Met	Asn	Ser	Glu	Arg	Pro	Asp	Glu	Asn	His	Ser	Glu	Pro	Ser	His
		755					760					765			
Gln	Asp	Val	He	Lys	Val	Lys	Glu	Glu	Phe	Thr	Asp	Pro	Thr	Tyr	Asp
	770					775					780				
Met	Phe	Tyr	Met	Ser	Gln	Tyr	Gly	Leu	Tyr	Asn	Gly	Gly	Gly	Ala	Ser
785					790					795					800
Met	Ala	Ala	Leu	His	Glu	Ser	Phe	Thr	Ser	Ser	Leu	Asn	Tyr	Gly	Ser
				805					810					815	
Pro	Gln	Lys	Phe	Ser	Pro	Glu	Gly	Asp	Leu	Cys	Ser	Ser	Pro	Asp	Pro
			820					825					830		
Lys	lle	Cys	Tyr	Val	Cys	Lys	Lys	Ser	Phe	Lys	Ser	Ser	Tyr	Ser	Val
		835					840					845			
Lys	Leu	His	Tyr	Arg	Asn	Val	His	Leu	Lys	Glu	Met	His	Val	Cys	Thr
	850					855					860				
Val	Ala	Gly	Cys	Asn	Ala	Ala	Phe	Pro	Ser	Arg	Arg	Ser	Arg	Asp	Arg
865					870					875					880
His	Ser	Ala	Asn	11e	Asn	Leu	His	Arg	Lys	Leu	Leu	Thr	Lys	Glu	Leu
				885					890					895	
Asp	Asp	Met	Gly	Leu	Asp	Ser	Ser	Gln	Pro	Ser	Leu	Ser	Lys	Asp	Leu
			900					905					910		
Arg	Asp	Glu	Phe	Leu	Val	Lys	lle	Tyr	Gly	Ala	Gln	His	Pro	Met	Gly
		915					920					925			
Leu	Asp	Val	Arg	Glu	Asp	Ala	Ser	Ser	Pro	Ala	Gly	Thr	Glu	Asp	Ser
	930					935					940				

His Leu Asn Gly Tyr Gly Arg Gly Met Ala Glu Asp Tyr Met Val Leu Asp Leu Ser Thr Thr Ser Ser Leu Gln Ser Ser Ser Ser Ile His Ser Ser Arg Glu Ser Asp Ala Gly Ser Asp Glu Gly Ile Leu Leu Asp Asp lle Asp Gly Ala Ser Asp Ser Gly Glu Ser Ala His Lys Ala Glu Ala Pro Ala Leu Pro Gly Ser Leu Gly Ala Glu Val Ser Gly Ser Leu Met Phe Ser Ser Leu Ser Gly Ser Asn Gly Gly Ile Met Cys Asn Ile Cys His Lys Met Tyr Ser Asn Lys Gly Thr Leu Arg Val His Tyr Lys Thr Val His Leu Arg Glu Met His Lys Cys Lys Val Pro Gly Cys Asn Met Met Phe Ser Ser Val Arg Ser Arg Asn Arg His Ser Gln Asn Pro Asn Leu His Lys Asn Ile Pro Phe Thr Ser Val Asp

<210> 4772

<211> 247

<212> PRT

<213> Homo sapiens

<400> 4772

Ala Ala Ala Leu Ala Leu Leu Thr Gly Gly Glu Met Leu Leu Asn Val Ala Leu Val Ala Leu Val Leu Leu Gly Ala Tyr Arg Leu Trp Val Arg Trp Gly Arg Gly Leu Gly Ala Gly Ala Gly Ala Gly Glu Glu Ser Pro Ala Thr Ser Leu Pro Arg Met Lys Lys Arg Asp Phe Ser Leu Glu Gln Leu Arg Gln Tyr Asp Gly Ser Arg Asn Pro Arg Ile Leu Leu Ala Val Asn Gly Lys Val Phe Asp Val Thr Lys Gly Ser Lys Phe Tyr Gly Pro Ala Gly Pro Tyr Gly lle Phe Ala Gly Arg Asp Ala Ser Arg Gly Leu Ala Thr Phe Cys Leu Asp Lys Asp Ala Leu Arg Asp Glu Tyr Asp Asp Leu Ser Asp Leu Asn Ala Val Gln Met Glu Ser Val Arg Glu Trp Glu Met Gln Phe Lys Glu Lys Tyr Asp Tyr Val Gly Arg Leu Leu Lys Pro Gly Glu Glu Pro Ser Glu Tyr Thr Asp Glu Glu Asp Thr Lys Asp His Asn Lys Gln Asp

<210> 4773

<211> 705

<212> PRT

<213> Homo sapiens

<400> 4773

Leu	Leu		Asp	Ser	Gln	Asp		Pro	Val	Thr	Val		Pro	Leu	Arg
		35					40					45			
Ala	Glu	Arg	Val	Leu	Leu	Phe	Asp	Asp	Ala	Leu		Leu	Leu	GIn	Gly
	50					55					60				
His	Asn	Val	His	Thr	Phe	Asp	Leu	Lys	Leu	Val	Trp	Val	Asp	Pro	
65					70					75					80
Gln	Asp	Gly	Cys	Thr	Phe	His	Leu	Leu	Thr	Pro	Glu	Glu	Glu	Phe	Ser
				85					90					95	
Phe	Cys	Ala	Lys	Asp	Ser	G1n	Gly	Gln	Ala	Val	Trp	Gln	Trp	Lys	Val
			100					105					110		
Thr	Trp	Ala	Val	His	Gln	Ala	Leu	His	Gly	Lys	Lys	Asp	Phe	Pro	Val
		115					120					125			
Leu	Gly	Ala	Gly	Leu	Glu	Pro	Ser	Gln	Pro	Pro	Asp	Cys	Arg	Cys	Ala
	130					135					140				
Glu	Tyr	Thr	Phe	Gln	Ala	Glu	Gly	Arg	Leu	Cys	Gln	Ala	Thr	Tyr	Glu
145					150					155					160
Gly	Glu	Trp	Cys	Arg	Gly	Arg	Pro	His	Gly	Lys	Gly	Thr	Leu	Lys	Trp
				165					170					175	
Pro	Asp	Gly	Arg	Asn	His	Val	Gly	Asn	Phe	Cys	Gln	Gly	Leu	Glu	His
			180					185					190		
Gly	Phe	Gly	He	Arg	Leu	Leu	Pro	Gln	Ala	Ser	Glu	Asp	Lys	Phe	Asp
		195					200					205			
Cys	Tyr	Lys	Cys	His	Trp	Arg	Glu	Gly	Ser	Met	Cys	Gly	Tyr	Gly	Пе
	210					215					220				
Cys	Glu	Tyr	Ser	Thr	Asp	Glu	Val	Tyr	Lys	Gly	Tyr	Phe	Gln	Glu	Gly
225					230					235					240
Leu	Arg	His	Gly	Phe	Gly	Val	Leu	Glu	Ser	Gly	Pro	Gln	Ala	Pro	Gln
				245					250					255	
Pro	Phe	Arg	Tyr	Thr	Gly	His	Trp	Glu	Arg	Gly	Gln	Arg	Ser	Gly	Tyr
			260					265					270		
Gly	lle	Glu	Glu	Asp	Gly	Asp	Arg	Gly	Glu	Arg	Tyr	lle	Gly	Met	Trp
		275					280					285			
Gln	Ala	Gly	Głn	Arg	His	G1 y	Pro	Gly	Val	Met	Val	Thr	Gln	Ala	Gly
	290					295					300				
Val	Cys	Tyr	Gln	Gly	Thr	Phe	G1n	Ala	Asp	Lys	Thr	Val	Gly	Pro	Gly
305					310					315					320

11e	Leu	Leu	Ser	Glu	Asp	Asp	Ser	Leu	lyr	61u	ыу	Inr	Phe	Inr	Arg
				325					330					335	
Asp	Leu	Thr	Leu	Met	Gly	Lys	Gly	Lys	Val	Thr	Phe	Pro	Asn	Gly	Phe
			340					345					350		
Thr	Leu	Glu	Gly	Ser	Phe	Gly	Ser	Gly	Ala	Gly	Arg	Gly	Leu	His	Thr
		355					360					365			
Gln	Gly	Val	Leu	Asp	Thr	Ala	Ala	Leu	Pro	Pro	Asp	Pro	Ser	Ser	Thr
	370					375					380				
Cys	Lys	Arg	Gln	Leu	Gly	Val	Gly	Ala	Phe	Pro	Val	Glu	Ser	Arg	Trp
385					390					395					400
G1n	Gly	Val	Tyr	Ser	Pro	Phe	Arg	Asp	Phe	Val	Cys	Ala	Gly	Cys	Pro
				405					410					415	
Arg	Asp	Leu	Gln	Glu	Ala	Leu	Leu	Gly	Phe	Asp	Val	Gln	Ser	Ser	Arg
			420					425					430		
Glu	Leu	Arg	Arg	Ser	Gln	Asp	Tyr	Leu	Ser	Cys	Glu	Arg	Thr	His	Pro
		435					440					445			
Glu	Asp	Ser	Val	Gly	Ser	Met	Glu	Asp	Ile	Leu	Glu	Glu	Leu	Leu	Gln
	450					455					460				
His	Arg	Glu	Pro	Lys	Ala	Leu	Gln	Leu	Tyr	Leu	Arg	Lys	Ala	Leu	Ser
465					470					475					480
Asn	Ser	Leu	His	Pro	Leu	Gly	Lys	Leu	Leu	Arg	Thr	Leu	Met	Leu	Thr
				485					490					495	
Phe	Gln	Ala	Thr	Tyr	Ala	Gly	Val	Gly	Ala	Asn	Lys	His	Leu	Gln	Gly
			500					505					510		
Leu	Ala	Gln	Glu	Glu	Val	Lys	Gln	His	Ala	Gln	Glu	Leu	Trp	Ala	Λla
		515					520					525			
Tyr	Arg	Gly	Leu	Leu	Arg	Val	Ala	Leu	Glu	Arg	Lys	Gly	Gln	Ala	Leu
	530					535					540				
Glu	Glu	Asp	Glu	Asp	Thr	Glu	Thr	Arg	Asp	Leu	Gln	Val	His	Gly	Leu
545					550					555					560
Val	Leu	Pro	Leu	Met	Leu	Pro	Ser	Phe	Tyr	Ser	Glu	Leu	Phe	Thr	Leu
				565					570					575	
Tyr	Leu	Leu	Leu	His	Glu	Arg	Glu	Asp	Ser	Phe	Tyr	Ser	Gln	Gly	He
			580					585					590		
Ala	Asn	Leu	Ser	Leu	Phe	Pro	Asp	Thr	Gln	Leu	Leu	Glu	Phe	Leu	Asp
		595					600					605			

Val Gln Lys His Leu Trp Pro Leu Lys Asp Leu Thr Leu Thr Ser Asn Gln Arg Tyr Ser Leu Val Arg Asp Lys Cys Phe Leu Ser Ala Thr Glu Cys Leu Gln Lys Ile Met Thr Thr Val Asp Pro Arg Glu Lys Leu Glu Val Leu Glu Arg Thr Tyr Gly Glu Ile Glu Gly Thr Val Ser Arg Val Leu Gly Arg Glu Tyr Lys Arg Pro Trp Thr Thr Cys Cys His Phe Ser Ser Thr Trp Cys Arg Ala Pro Glu Phe Ser Thr Trp Glu Pro Arg Ser Thr <210> 4774 <211> 907 <212> PRT <213> Homo sapiens <400> 4774 Met Gln Tyr Gln Cys Lys Lys Cys Asn Val Val Phe Pro Arg Ile Phe Asp Leu Ile Thr His Gln Lys Lys Gln Cys Tyr Lys Asp Glu Asp Asp Asp Ala Gln Asp Glu Ser Gln Thr Glu Asp Ser Met Asp Ala Thr Asp Gln Val Val Tyr Lys His Cys Thr Val Ser Gly Gln Thr Asp Ala Ala Lys Asn Ala Ala Ala Pro Ala Ala Ser Ser Gly Ser Gly Thr Ser Thr Pro Leu IIe Pro Ser Pro Lys Pro Glu Pro Glu Lys Thr Ser Pro Lys Pro Glu Tyr Pro Ala Glu Lys Pro Lys Gln Ser Asp Pro Ser Pro Pro

Ser	Gln	Gly	Thr	Lys	Pro	Ala	Leu	Pro	Leu	Ala	Ser	Thr	Ser	Ser	Asp
		115					120					125			
Pro	Pro	Gln	Ala	Ser	Thr	Ala	Gln	Pro	G1n	Pro	Gln	Pro	GIn	Pro	Pro
	130					135					140				
Lys	Gln	Pro	Gln	Leu	Пе	Gly	Arg	Pro	Pro	Ser	Ala	Ser	Gln	Thr	Pro
145					150					155					160
Val	Pro	Ser	Ser	Pro	Leu	Gln	Пе	Ser	Met	Thr	Ser	Leu	Gln	Asn	Ser
				165					170					175	
Leu	Pro	Pro	Gln	Leu	Leu	Gln	Tyr	Gln	Cys	Asp	Gln	Cys	Thr	Val	Ala
			180					185					190		
Phe	Pro	Thr	Leu	Glu	Leu	Trp	Gln	Glu	His	Gln	His	Val	His	Phe	Leu
		195					200					205			
Ala	Ala	G1n	Asn	GIn	Phe	Leu	His	Ser	Pro	Phe	Leu	Glu	Arg	Pro	Met
	210					215					220				
Asp	Met	Pro	Tyr	Met	He	Phe	Asp	Pro	Asn	Asn	Pro	Leu	Met	Thr	Gly
225					230					235					240
Gln	Leu	Leu	Gly	Ser	Ser	Leu	Thr	Gln	Met	Pro	Pro	Gln	Ala	Ser	Ser
				245					250					255	
Ser	His	Thr	Thr	Ala	Pro	Thr	Thr	Val	Ala	Ala	Ser	Leu	Lys	Arg	Lys
			260					265					270		
Leu	Asp	Asp	Lys	Glu	Asp	Asn	Asn	Cys	Ser	Glu	Lys	Glu	G1 y	Gly	Asn
		275					280		•			285			
Ser	Gly	Glu	Asp	Gln	His	Arg	Asp	Lys	Arg	Leu	Arg	Thr	Thr	He	Thr
	290					295					300				
Pro	Glu	Gln	Leu	Glu	He	Leu	Tyr	Glu	Lys	Tyr	Leu	Leu	Asp	Ser	Asn
305					310					315					320
Pro	Thr	Arg	Lys	Met	Leu	Asp	His	lle	Ala	Arg	Glu	Val	G1 y	Leu	Lys
				325					330					335	
Lys	Arg	Val	Val	Gln	Val	Trp	Phe	Gln	Asn	Thr	Arg	Ala	Arg	Glu	Arg
			340					345					350		
Lys	Gly	Gln	Phe	Arg	Ala	Val	G]y	Pro	Ala	Gln	Ser	His	Lys	Arg	Cys
		355					360					365			
Pro	Phe	Cys	Arg	Ala	Leu	Phe	Lys	Ala	Lys	Ser	Ala	Leu	Glu	Ser	His
	370					375					380				

lle Arg Ser Arg His Trp Asn Glu Gly Lys Gln Ala Gly Tyr Ser Leu

385					390					395					400
Pro	Pro	Ser	Pro	Leu	Ile	Ser	Thr	Glu	Asp	Gly	Gly	Glu	Ser	Pro	G1n
				405					410					415	
Lys	Tyr	He	Tyr	Phe	Asp	Tyr	Pro	Ser	Leu	Pro	Leu	Thr	Lys	11e	Asp
			420					425					430		
Leu	Ser	Ser	Glu	Asn	Glu	Leu	Ala	Ser	Thr	Va]	Ser	Thr	Pro	Val	Ser
		435					440					445			
Lys	Thr	Ala	Glu	Leu	Ser	Pro	Lys	Asn	Leu	Leu	Ser	Pro	Ser	Ser	Phe
	450					455					460				
Lys	Ala	Glu	Cys	Ser	Glu	Asp	Val	Glu	Asn	Leu	Asn	Ala	Pro	Pro	Ala
465					470					475					480
Glu	Ala	Gly	Tyr	Asp	Gln	Asn	Lys	Thr	Asp	Phe	Asp	Glu	Thr	Ser	Ser
				485					490					495	
He	Asn	Thr	Ala	Ile	Ser	Asp	Ala	Thr	Thr	Gly	Asp	Glu	Gly	Așn	Thr
			500					505					510		
Glu	Met	Glu	Ser	Thr	Thr	Gly	Ser	Ser	Gly	Asp	Val	Lys	Pro	Ala	Leu
		515					520					525			
Ser	Pro	Lys	Glu	Pro	Lys	Thr	Leu	Asp	Thr	Leu	Pro	Lys	Pro	Ala	Thr
	530					535					540				
Thr	Pro	Thr	Thr	Glu	Val	Cys	Asp	Asp	Lys	Phe	Leu	Phe	Ser	Leu	Thr
545					550					555					560
Ser	Pro	Ser	He	His	Phe	Asn	Asp	Lys	Asp	G1 y	Asp	His	Asp	Gln	Ser
				565					570					575	
Phe	Tyr	lle	Thr	Asp	Asp	Pro	Asp	Asp	Asn	Ala	Asp	Arg	Gly	Glu	Thr
			580					585					590		
Ser	Ser	He	Ala	Asp	Pro	Ser	Ser	Pro	Asn	Pro	Phe	Gly	Ser	Ser	Asn
		595					600					605			
Pro		Lys	Ser	Lys	Ser		Asp	Arg	Pro	G1 y		Lys	Arg	Phe	Arg
	610					615					620				
Thr	Gln	Met	Ser	Asn		G1n	Leu	Lys	Val		Lys	Ala	Cys	Phe	
625					630					635					640
Asp	Tyr	Arg	Thr		Thr	Met	Gln	Glu	-	Glu	Met	Leu	Gly		Glu
				645					650					655	
He	Gly	Leu		Lys	Arg	Val	Val		Val	Trp	Phe	Gln	Asn	Ala	Arg
			660				20.	665				_	670		. .
110	Lvc	Chi	1 22 0	Lvc	Pho	WC	110	Acn	110	G1v	LVC	Pro	Pho	Met	Ha

		675					680					685			
Asn	Gln	Gly	Gly	Thr	Glu	Gly	Thr	Lys	Pro	Glu	Cys	Thr	Leu	Cys	Gly
	690					695					700				
Val	Lys	Tyr	Ser	Λla	Arg	Leu	Ser	11e	Arg	Asp	His	Пe	Phe	Ser	Lys
705					710					715					720
Gln	His	lle	Ser	Lys	Val	Arg	Glu	Thr	Val	Gly	Ser	Gln	Leu	Asp	Arg
				725					730					735	
Glu	Lys	Asp	Tyr	Leu	Ala	Pro	Thr	Thr	Val	Arg	Gln	Leu	Met	Ala	Gln
			740					745					750		
Gln	Glu	Leu	Asp	Arg	Ile	Lys	Lys	Ala	Ser	Asp	Val	Leu	Gly	Leu	Thr
		755					760					765			
Val	Gln	Gln	Pro	Gly	Met	Met	Asp	Ser	Ser	Ser	Leu	His	Gly	lle	Ser
	770					775					780				
Leu	Pro	Thr	Λla	Tyr	Pro	Gly	Leu	Pro	Gly	Leu	Pro	Pro	Val	Leu	Leu
785					790					795					800
Pro	Gly	Met	Asn	G1 y	Pro	Ser	Ser	Leu	Pro	Gly	Phe	Pro	G1n	Asn	Ser
				805					810					815	
Asn	Thr	Leu	Thr	Pro	Pro	Gly	Ala	Gly	Met	Leu	Gly	Phe	Pro	Thr	Ser
			820					825					830		
Ala	Thr	Ser	Ser	Pro	Ala	Leu	Ser	Leu	Ser	Ser	Ala	Pro	Thr	Lys	Pro
		835					840					845			
Leu	Leu	Gln	Thr	Pro	Pro	Pro	Pro	Pro	Pro	Pro	Pro	Pro	Pro	Pro	Pro
	850					855					860				
Ser	Ser	Ser	Leu	Ser	Gly	Gln	Gln	Thr	Glu	Gln	Gln	Asn	Lys	Glu	Ser
865					870					875					880
Glu	Lys	Lys	Gln	Thr	Lys	Pro	Asn	Lys	Val	Lys	Lys	He	Lys	Glu	Glu
				885					890					895	
Glu	Leu	Glu	Ala	Thr	Lys	Pro	Glu	Lys	His	Pro					
			900					905							

<211> 412

<212> PRT

<213> Homo sapiens

<400)> 47	775													
Met	Thr	Phe	Thr	Phe	Gln	Ser	Glu	Asp	Leu	Lys	Arg	Asp	Cys	Gly	Lys
1				5					10					15	
Lys	Met	Ser	His	Gln	His	Val	Phe	Ser	Leu	Ala	Met	Glu	Glu	Asp	Val
			20					25					30		
Lys	Thr	Ala	Asp	Thr	Lys	Lys	Ala	Ser	Arg	lle	Leu	Asp	His	Glu	Lys
		35					40					45			
Glu	Asn	Thr	Arg	Ser	lle	Cys	Leu	Leu	Glu	Gln	Lys	Arg	Lys	Val	Val
	50					55					60				
Ser	Ser	Asn	Ile	Asp	Val	Pro	Pro	Ala	Arg	Lys	Ser	Ser	Glu	Glu	Leu
65					70					75					80
Asp	Met	Asp	Lys	Val	Thr	Ala	Ala	Met	Val	Leu	Thr	Ser	Leu	Ser	Thr
				85					90					95	
Ser	Pro	Leu	Val	Arg	Ser	Pro	Pro	Val	Arg	Pro	Asn	Glu	Ser	Leu	Ser
			100					105					110		
Gly	Ser	Trp	Lys	Glu	G1 y	Gly	Cys	Val	Pro	Ser	Ser	Thr	Ser	Ser	Ser
		115					120					125			
Gly	Tyr	Trp	Ser	Trp	Ser	Ala	Pro	Ser	Asp	Gln	Ser	Asn	Pro	Ser	Thr
	130					135					140				
Pro	Ser	Pro	Pro	Leu	Ser	Ala	Asp	Ser	Phe	Lys	Pro	Phe	Arg	Ser	Pro
145					150					155					160
Ala	Gln	Pro	Asp	Asp	Gly	Ile	Asp	G] u	Ala	G] u	Ala	Ser	Asn	Leu	Leu
				165					170					175	
Phe	Asp	Glu		He	Pro	Arg	Lys		Lys	Asn	Ser	Met		Val	Met
			180					185					190		
Phe	Lys		Leu	Trp	Lys	Asn		Gly	Lys	Val	Leu		Thr	Ala	Ala
63	7.3	195	,		1.7		200	7.1			61	205	17 7	<i>(</i>)	
Gly		Gln	Lys	His	He	Arg	Thr	He	His	Leu		Arg	Val	Gly	Asp
C	210	т	C	Δ.	C 1	215	C1		DI	т.	220	TL	C1	7 1	1
	Asp	ıyr	5er	Asp		Glu	61u	Asp	rne		lyr	ınr	Glu	11e	
225		TI		C	230	A 3		C1	1	235	C		4.7	15	240
Leu	Asn	Ihr	Asp		val	Ala	Asp	Gly		Ser	Ser	Leu	Ala		val
6	n.	C	C1	245		A 3	C	n.	250 D	TI	DI	D.	77	255 D	Δ
Ser	Pro	Ser		Ser	Leu	Ala	5er		Pro	Ihr	Phe	Pro		Pro	Asp
C = -	C =	۸	260	C1.	Tl	D	C	265	1	Tl	C1.	Tl	270	1	M = *
5er	261.	лrg	ınr	61u	inr	Pro	Cys	Ala	Lys	ınr	GIU	ınr	Lys	Leu	met

Thr Pro Leu Ser Arg Ser Ala Pro Thr Thr Leu Tyr Leu Val His Thr Asp His Ala Tyr Gln Ala Thr Pro Pro Val Thr Ile Pro Gly Ser Ala Lys Phe Thr Pro Asn Gly Ser Ser Phe Ser Ile Ser Trp Gln Ser Pro Pro Val Thr Phe Thr Gly Ile Pro Val Ser Pro Thr His His Pro Val Gly Thr Gly Glu Gln Arg Gln His Ala His Thr Val Leu Ser Ser Pro Pro Arg Gly Thr Val Ser Leu Arg Lys Pro Arg Gly Glu Gly Lys Lys Cys Arg Lys Val Tyr Gly Met Glu Asn Arg Asp Met Trp Cys Thr Ala Cys Arg Trp Lys Lys Ala Cys Gln Arg Phe Leu Asp

<210> 4776

<211> 314

<212> PRT

<213> Homo sapiens

<400> 4776

Lys Thr Asp Phe Lys Pro Thr Lys Ile Lys Arg Asp Lys Glu Gly His

				85					90					95	
Tyr	Thr	Met	Val	Lys	Gly	Ser	He	Gln	Gln	Glu	Glu	Leu	Thr	Πe	Leu
			100					105					110		
Asn	He	Tyr	Ala	Pro	Asn	Thr	Gly	Ala	Pro	Arg	Phe	He	Lys	G1n	Val
		115					120					125			
Leu	Ser	Asp	Val	Gln	Arg	Asp	Leu	Asp	Ser	His	Thr	Leu	Ile	Met	Gly
	130					135					140				
Asp	Phe	Asn	Thr	Pro	Leu	Ser	Thr	Leu	Asp	Arg	Ser	Thr	Arg	Gln	Lys
145					150					155					160
Val	Asn	Lys	His	Thr	Gln	Glu	Phe	Asn	Ser	Ala	Leu	His	G1n	Ala	Asp
				165					170					175	
Leu	Ile	Asp	He	Tyr	Lys	Thr	Leu	His	Pro	Lys	Ser	Thr	Glu	Tyr	Thr
			180					185					190		
Phe	Phe	Ser	Ala	Pro	His	His	Thr	Tyr	Ser	Lys	He	Asp	His	Ile	Phe
		195					200					205			
Gly	Ser	Lys	Ala	Leu	Phe	Ser	Lys	Cys	Lys	Arg	Thr	Glu	He	He	Thr
	210					215					220				
Asn	Cys	Leu	Ser	Glu	His	Ser	Ala	lle	Lys	Leu	Glu	Leu	Arg	Ile	Lys
225					230					235					240
Asn	Leu	Thr	Gln	Asn	Arg	Ser	Thr	Thr	Trp	Lys	Leu	Asn	Asn	Leu	Leu
				245					250					255	
Leu	Asn	Asp	Tyr	Trp	Val	His	Asn	Glu	Met	Lys	Ala	Glu	Ile	Lys	Met
			260					265					270		
Phe	Phe	Glu	Thr	Asn	Glu	Asn	Lys	Asp	Thr	Thr	Tyr	Gln	Asn	Leu	Trp
		275					280					285			
Asp	Ala	Phe	Lys	Ala	Val	Cys	Thr	Gly	Lys	Phe	Ile	Ala	Leu	Asn	Ala
	290					295					300				
His	Lys	Arg	Lys	Lys	Glu	Arg	Lys	Lys	Glu						
305					310										

<211> 264

<212> PRT

<213> Homo sapiens

<400)> 47	777													
Met	Thr	G1 y	Ser	Λsn	Ser	His	Ile	Thr	Ile	Leu	Thr	Leu	Asn	Val	Asn
1				5					10					15	
Gly	Leu	Asn	Thr	Pro	lle	Lys	Arg	His	Arg	Leu	Ala	Ser	Trp	lle	Lys
			20					25					30		
Ser	Gln	Asp		Ser	Val	Cys	Cvs	Ile	Gln	Glu	Thr	His	Leu	He	Cvs
		35				-,-	40					45			•
lve	Asn		Tyr	Aro	Len	Lys		lvs	Glv	Trp	Arø		He	Tvr	Gln
Lys	50	Mid	1 9 1	ni 8	Lcu	55	110	Lys	019	II p	60	LJO	110	.,1	01 111
Ala		A 200	Luc	Cln	Luc	Lys	A10	Cly	Vol.	41 0		Lou	Va1	Sor	Acn
	ASII	AI g	Lys	0111		Lys	пта	Oly	val		116	Leu	vai	561	80
65	C		DI.	,	70 D	TI.		11.1	1	75	A	1	1	C1	
Lys	Ser	Asp	Phe		ro	Thr	Lys	11e		Arg	Asp	Lys	Lys		піѕ
_				85			.1	0.1	90	61	0.3		mı.	95	
Tyr	He	Met		Lys	Gly	Ser	Met		GIn	Glu	Glu	Leu		He	Leu
			100					105					110		
Asn	He	Tyr	Ala	Pro	Asn	Thr	G1 y	Ala	Pro	Arg	Phe	Ile	Asn	Glu	Val
		115					120					125			
Leu	Arg	Asp	Pro	Gln	Arg	Asn	Leu	Asp	Ser	His	Thr	lle	Ile	Met	Gly
	130					135					140				
Asp	Phe	Asn	Thr	Pro	Leu	Ser	lle	Phe	Asp	Arg	Ser	Thr	Arg	Gln	Lys
145					150					155					160
Val	Asn	Lys	Asp	Thr	Gln	Asp	Leu	Asn	Leu	Thr	Leu	Gln	Gln	Ala	Asp
				165					170					175	
Leu	Arg	Asp	He	Tyr	Arg	Thr	Leu	His	Pro	Lys	Ser	Thr	Glu	Tyr	Thr
			180					185					190		
Phe	Phe	Ser	Ala	Leu	His	His	Thr	Tyr	Ser	Lys	Thr	Asp	His	He	Leu
		195					200					205			
Gly	Ser	Lys	Ala	Leu	Leu	Ser	Lys	Cys	Lys	Arg	Thr	Glu	lle	Thr	Thr
	210	·				215					220				
Asn		Leu	Ser	Asp	His	Ser	Ala	lle	Lys	Leu		Leu	Arg	Ile	Lys
225	-, -			- 1-	230				, -	235		-			240
	l eu	Thr	Gln	Tvr		lle	Ser	Thr	Trn		l.eu	Asn	Asn	Leu	
L) 3	1,0 u	1111	0111	245	1110	110	001		250	22,5	nea	,,,,,,,		255	.,,,,
l en	Aen	Asn	Tvr		Val	Asn	Asn		200						
Leu	11011	nap	ı yı	ıтЫ	101	11011	11011								

```
<210> 4778
〈211〉 852
<212> PRT
<213> Homo sapiens
<400> 4778
Met Leu Ile Thr Glu Arg Lys His Phe Arg Ser Gly Arg Ile Ala Gln
                                      10
 1
Ser Met Ser Glu Ala Asn Leu Ile Asp Met Glu Ala Gly Lys Val Ser
Lys Ser Cys Asn Ile Thr Glu Cys Gln Asp Pro Asp Leu Leu His Asn
         35
                             40
                                                  45
Trp Pro Asp Ala Phe Thr Leu Arg Gly Asn Asn Ala Ser Lys Val Ala
     50
                         55
                                              60
Asn Pro Phe Trp Asn Gln Leu Ser Ala Ser Asn Pro Phe Leu Asp Asp
                     70
                                          75
Ile Thr Gln Leu Arg Asn Asn Arg Lys Arg Asn Asn Ile Ser Ile Leu
                                      90
                 85
Lys Glu Asp Pro Phe Leu Phe Cys Arg Glu Ile Glu Asn Gly Asn Ser
                                 105
Phe Asp Ser Ser Gly Asp Glu Leu Asp Val His Gln Leu Leu Arg Gln
        115
                                                 125
                            120
Thr Ser Ser Arg Asn Ser Gly Arg Ser Lys Ser Val Ser Glu Leu Leu
    130
                        135
                                             140
Asp Ile Leu Asp Asp Thr Ala His Ala His Gln Ser Ile His Asn Ser
                    150
                                         155
Asp Gln 11e Leu Leu His Asp Leu Glu Trp Pro Lys Asn Asp Arg Glu
                165
Ala Tyr Lys Met Ala Trp Leu Ser Gln Arg Gln Leu Ala Arg Ser Cys
                                 185
Leu Asp Leu Asn Thr Ile Ser Gln Ser Pro Gly Trp Ala Gln Thr Gln
                            200
                                                 205
        195
Leu Ala Glu Val Thr lle Ala Cys Lys Val Asn His Gln Gly Gly Ser
```

Val	Gln	Leu	Pro	Glu	Ser	Asp	He	Thr	Val	His	Val	Pro	Gln	Gly	His
225					230					235					240
Val	Ala	Val	Gly	Glu	Phe	Gln	Glu	Val	Ser	Leu	Arg	Ala	Phe	Leu	Asp
				245					250					255	
Pro	Pro	His	Met	Leu	Asn	His	Asn	Leu	Ser	Cys	Thr	Val	Ser	Pro	Leu
			260					265					270		
Leu	Glu	Ile	Met	Leu	Gly	Asn	Leu	Asn	Thr	Met	Glu	Ala	Leu	Leu	Leu
		275					280					285			
Glu	Met	Lys	Ile	Gly	Ala	Glu	Val	Arg	Lys	Asp	Pro	Phe	Ser	Gln	Val
	290					295					300				
Met	Thr	Glu	Met	Val	Cys	Leu	His	Ser	Leu	Gly	Lys	Glu	G1 y	Pro	Phe
305					310					315					320
Lys	Val	Leu	Ser	Asn	Cys	Tyr	lle	Tyr	Lys	Asp	Thr	lle	Gln	Val	Lys
				325					330					335	
Leu	lle	Asp	Leu	Ser	GIn	Val	Met	Tyr	Leu	Val	Val	Ala	Ala	Gln	Ala
			340					345					350		
Lys	Ala	Leu	Pro	Ser	Pro	Ala	Ala	Thr	He	Trp	Asp	Tyr	Ile	His	Lys
		355					360					365			
Thr	Thr	Ser	lle	Gly	lle	Tyr	Gly	Pro	Lys	Tyr	lle	His	Pro	Asn	Phe
	370					375					380				
Thr	Val	Val	Leu	Thr	Val	Cys	Gly	His	Asn	Tyr	Met	Pro	Gly	Gln	Leu
385					390					395					400
Thr	He	Ser	Asp	lle	Lys	Lys	Gly	Gly	Lys	Asn	He	Ser	Pro	Val	Val
				405					410					415	
Phe	Gln	Leu	Trp	Gly	Lys	Gln	Ser	Phe	Leu	Leu	Asp	Lys	Pro	Gln	Asp
			420					425					430		
Leu	Ser	He	Ser	He	Phe	Ser	Cys	Asp	Pro	Asp	Phe	Glu	Val	Lys	Thr
		435					440					445			
G]u	Gly	Glu	Arg	Lys	Glu	He	Lys	Gln	Lys	Gln	Leu	Glu	Ala	Gly	Glu
	450					455					460				
Val	Val	His	Gln	Gln	Phe	Leu	Phe	Ser	Leu	Val	Glu	His	Arg	Glu	Met
465					470					475					480
His	Leu	Phe	Asp	Phe	Cys	Val	Gln	Val	Glu	Pro	Pro	Asn	Gly	Glu	Pro
				485					490					495	
Val	Ala	Gln	Phe	Ser	lle	Thr	Thr	Pro	Asp	Pro	Thr	Pro	Asn	Leu	Lys
			500					505					510		

Arg	Leu	Leu	Asn	Leu	Pro	Gly	Tyr	Leu	Gln	Lys	Lys	Glu	Glu	lle	Lys
		515					520					525			
Ser	Ala	Pro	Leu	Ser	Pro	Lys	He	Leu	Val	Lys	Tyr	Pro	Thr	Phe	Gln
	530					535					540				
Asp	Lys	Thr	Leu	Asn	Phe	Ser	Asn	Tyr	G1 y	Val	Thr	Leu	Lys	Ala	Val
545					550					555					560
Leu	Arg	Gln	Ser	Lys	lle	Asp	Tyr	Phe	Leu	Glu	Tyr	Phe	Lys	G1 y	Asp
				565					570					575	
Thr	He	Ala		Leu	Gly	Glu	Gly		Val	Lys	Ala	Ile		G1n	Ser
			580					585					590		
Lys	Val		Glu	Trp	Tyr	Val		Val	Leu	Arg	Gly		He	Gly	Leu
		595					600					605			
Val		Cys	Lys	Asn	Va]	Lys	Val	He	Ser	Lys		Gln	Val	Met	Phe
	610		_			615					620				
	Ser	Asp	Ser	Val		Thr	Thr	Arg	Asn		Leu	Glu	GIn	He	
625	_				630					635				6 01	640
Leu	Pro	Leu	Lys		Leu	Thr	Tyr	He		Ser	Val	Val	Leu		Leu
V 3	c	61		645	т		т		650	,	4.1		V 7	655	61
Val	Ser	GJu		Val	lyr	Asp	1rp		vai	Leu	Ala	Asp		Leu	Gly
Т	C	u: a	660	C	1	C1.,	A ===	665	A a.s.	C1»	710	C1 m	670	A an	Lua
ıyr	ser		Leu	Ser	Leu	Glu		rne	ASP	GIN	11e		Ala	ASP	Lys
C1	Can	675	Luc	Vol	Con	Tun	680	110	Luc	Luc	Lau	685	C1	Aon	Cva
Giu	690	Giu	Lys	vai	361	Tyr 695	vai	116	Lys	Lys	700	Lys	Glu	nsp	Cys
Hic		Glu	Ara	Acn	The	Arg	Lvc	Pho	Lou	Tur		Lou	110	Val	ΔΊα
705	1111	Olu	Mig	11.511	710	Mg	1.73	1116	LCu	715	Olu	Lea	.110	, 01	720
	Leu	Lvs	Met	Asn		Gln	Gln	Leu	Val		Arg	Len	Tle	Gln	
1200	Lou	2,0		725	0,0	0111	010	Loc	730	1110	6	200		735	0.4
Ala	Ala	Val	Leu		Ser	Ala	Val	Lvs		Glv	Lvs	Glv	Trp		Glu
			740					745		,	, -		750	0	
Leu	Ala	Glu		Leu	Val	Arg	Leu		Lys	Gln	Gln	Met		Ala	Tyr
		755	•			ζ,	760		•			765			·
Glu	He	Pro	His	Arg	Gly	Asn	Thr	Gly	Asp	Val	Ala	Va]	Glu	Met	Met
	770			_	•	775		٠	•		780				
Trp	Lys	Pro	Ala	Tyr	Asp	Phe	Leu	Tyr	Thr	Trp	Ser	Ala	His	Tyr	Gly
785					790					795					800

<210> 4779

<211> 865

<212> PRT

<213> Homo sapiens

<400> 4779

 Met Ser Leu Gln Asp Ser Thr Leu Ser Arg Glu Gly Lys Pro Glu Gly

 1
 5
 10
 15

 Glu 1le Met Ala Ala Val Phe Phe Ser Val Gly Arg Leu Ser Pro Glu
 20
 25
 30

 Val Thr Gln Pro Asp Glu Asp Leu His Leu Gln Ala Glu Glu Thr Gln
 35
 40
 45

Leu Val Lys Glu Ser Val Thr Phe Lys Asp Val Ala Ile Asp Phe Thr

50 55 60

Leu Glu Glu Trp Arg Leu Met Asp Pro Thr Gln Arg Asn Leu His Lys
65 70 75 80

Asp Val Met Leu Glu Asn Tyr Arg Asn Leu Val Ser Leu Gly Leu Ala 85 90 95

Val Ser Lys Pro Asp Met Ile Ser His Leu Glu Asn Gly Lys Gly Pro 100 105 110

Trp Val Thr Val Arg Glu lle Ser Arg lle Pro Tyr Pro Asp Met Glu 115 120 125

Pro Lys Pro Ala Thr Lys Asn Ala Thr Arg Thr Lys Ala Ile Ser Glu 130 135 140

Asp Leu Ser Gln Glu Ala lle Leu Glu Lys Leu Thr Glu Asn Gly Leu 145 150 155 160

Trp	Asp	Ser	Arg	Met	Glu	Gly	Leu	Trp	Lys	Trp	Asn	Asp	Arg	Ile	Leu
				165					170					175	
Arg	Leu	Gln	Asn	Asn	Gln	Glu	Asn	His	Leu	Ser	Gln	Arg	Ile	lle	Pro
			180					185					190		
Leu	Lys	Lys	Thr	Pro	Thr	Ser	Gln	Arg	G1y	Phe	Arg	Phe	Glu	Ser	He
		195					200					205			
Leu	He	Pro	Glu	Pro	Gly	lle	Λla	Thr	Glu	Glu	Leu	His	Ser	Arg	Cys
	210					215					220				
Gln	Thr	Gln	Glu	Glu	Asn	Phe	Thr	Glu	Asn	Leu	Asn	Leu	Ile	Thr	Asp
225					230					235					240
Thr	His	Leu	Gly	Lys	lle	lle	Cys	Lys	Glu	Met	Lys	Gly	Ser	Lys	Ala
				245					250					255	
11e	Arg	Gln	Thr	Ser	Glu	Leu	Thr	Leu	Gly	Lys	Lys	Ser	Asn	Asn	Lys
			260					265					270		
Glu	Lys	Pro	Tyr	Lys	Cys	Ser	Thr	Cys	Glu	Lys	Ala	Phe	His	Tyr	Arg
		275					280					285			
Ser	Leu	Leu	Ile	Gln	His	Gln	Arg	Thr	His	Thr	Lys	Glu	Lys	Pro	Tyr
	290					295					300				
Glu	Cys	Asn	Glu	Cys	Gly	Lys	Thr	Phe	Ser	Gln	Pro	Ser	Tyr	Leu	Ser
305					310					315					320
Gln	His	Lys	Lys	He	His	Thr	Gly	Glu	Lys	Pro	Tyr	Lys	Cys	Asn	Glu
				325					330					335	
Cys	Gly	Lys	Ala	Phe	He	Ala	Ser	Ser	Ser	Leu	Met	Val	His	Gln	Arg
			340					345					350		
lle	His	Thr	Lys	Glu	Lys	Pro	Tyr	Gln	Cys	Asn	Val	Cys	Gly	Lys	Ser
		355					360					365			
Phe	Ser	Gln	Cys	Ala	Arg	Leu	Asn	Gln	His	Gln	Arg	He	Gln	Thr	Gly
	370					375					380				
Glu	Lys	Pro	Tyr	Lys	Cys	Ser	6] u	Cys	Gly	Lys	Ala	Phe	Ser	Asp	Lys
385					390					395					400
Ser	Lys	Leu	Ala	Arg	His	Gln	Glu	Thr	His	Asn	Gly	G]u	Lys	Pro	Tyr
				405					410					415	
Lys	Cys	Asp	Asp	Cys	Gly	Lys	Ala	Phe	Arg	Asn	Lys	Ser	Tyr	Leu	Ser
			420					425					430		
Val	His	Gln	Lys	Thr	His	Thr	Glu	Glu	Lys	Pro	Tyr	Gln	Cys	Asn	Glu
		435					440					445			

Cys	Gly	Lys	Ser	Phe	Lys	Asn	Thr	Thr	Ile	Phe	Asn	Val	His	Gln	Arg
	450					455					460				
He	His	Thr	Gly	Glu	Lys	Pro	Phe	Arg	Cys	Asn	Glu	Cys	Gly	Lys	Λla
465					470					475					480
Tyr	Arg	Ser	Asn	Ser	Ser	Leu	Пe	Val	His	11e	Arg	Thr	His	Thr	Gly
				485					490					495	
Glu	Lys	Pro	Tyr	Glu	Cys	Asn	Glu	Cys	Gly	Lys	Ala	Phe	Asn	Arg	He
			500					505					510		
Ala	Asn	Phe	Thr	Glu	His	Gln	Arg	Ile	His	Thr	Gly	Glu	Lys	Pro	Tyr
		515					520					525			
Lys	Cys	Asn	Glu	Cys	Gly	Lys	Ala	Phe	Πle	Asn	Tyr	Ser	Cys	Leu	Thr
	530					535					540				
Val	His	His	Arg	Met	His	Thr	Gly	Glu	Lys	Pro	Tyr	Lys	Cys	Thr	Glu
545					550					555					560
Cys	Gly	Lys	Ala	Phe	Met	Arg	Ser	Ser	Ser	Leu	He	He	His	Gln	Cys
				565					570					575	
Ile	His	Thr	Glu	Glu	Lys	Pro	Tyr	Leu	Cys	Asn	Glu	Cys	Gly	Glu	Ser
			580					585					590		
Phe	Arg	Ile	Lys	Ser	His	Leu	Thr	Val	His	Gln	Arg	He	His	Thr	Gly
		595					600					605			
Glu	Lys	Pro	Tyr	Lys	Cys	Thr	Asp	Cys	Glu	Arg	Ala	Phe	Thr	Lys	Met
	610					615					620				
Val	Asn	Leu	Lys	Glu	His	G1n	Lys	He	His	Thr	Gly	Val	Lys	Pro	Tyr
625					630					635					640
Lys	Cys	Tyr	Asp	Cys	Gly	Lys	Ser	Phe	Arg	Thr	Lys	Ser	Tyr	Leu	lle
				645					650					655	
Val	His	Gln	Arg	Thr	His	Thr	Gly	Glu	Lys	Pro	Tyr	Lys	Cys	Asn	Glu
			660					665					670		
Cys	Glu	Lys	Ala	Phe	Thr	Asn	Thr	Ser	Gln	Leu	Thr	Val	His	Gln	Arg
		675					680					685			
Arg	His	Thr	Gly	Glu	Lys	Pro	Tyr	Lys	Cys	Asn	Glu	Cys	Gly	Lys	Val
	690					695					700				
Phe	Thr	Ser	Asn	Ser	Gly	Phe	Asn	Thr	His	Gln	Arg	Thr	His	Thr	G1y
705					710					715					720
Glu	Lys	Pro	Phe		Cys	Asn	Asp	Cys	Gly	Lys	Ala	Phe	Ser	Gln	Met
				725					730					735	

```
Val His Val Thr Glu His Gln Lys Ile His Ser Gly Glu Lys Pro Tyr
            740
                                745
Lys Cys Asp Val Cys Gly Lys Ala Phe Arg Arg Gly Ser Tyr Leu Thr
        755
                            760
                                                 765
Val His Trp Arg Thr His Thr Gly Glu Lys Pro Tyr Thr Cys Lys Glu
                                             780
                        775
Cys Gly Lys Gly Cys Ile Thr Leu Ser Gln Leu Thr Leu His Gln Arg
                    790
                                        795
Ile His Thr Gly Glu Arg Pro Tyr Lys Cys Glu Glu Cys Gly Lys Ala
                                     810
                805
Phe Arg Thr Asn Ser Asp Phe Thr Val His Leu Arg Met His Thr Gly
                                825
Glu Lys Pro Tyr Lys Cys Asn Glu Cys Gly Lys Ala Phe Arg Ser Ser
        835
                            840
                                                 845
Ser Ser Leu Thr Val His Gln Arg Ile His Gln Arg Glu Thr Gln Leu
    850
                        855
                                             860
lle
865
<210> 4780
<211> 547
<212> PRT
<213> Homo sapiens
<400> 4780
Met Leu Glu Gly Arg Gln Thr Pro Ala Ser Thr Leu Glu Gln Asp Ala
                  5
                                     10
Thr Asp Tyr Arg Leu Arg Ser Leu Arg Lys Leu Leu Ala Gln Pro Arg
             20
                                 25
                                                      30
Glu Gly Leu Leu Ala Pro Phe Ser Lys Arg Asn Ser Thr Ala Ser Phe
                                                  45
```

Pro Gly Arg Thr Ser His 11e Pro Val Gln Gln Pro Glu Lys Arg Lys

Gln Lys Pro Ser Pro Glu Pro Ser Gln Asp Ser Pro His Ser Asp Lys

Trp	Pro	Pro	Gly	His	Pro	Val	Lys	Asn	Leu	Pro	Gln	Met	Arg	Gly	Pro
				85					90					95	
Arg	Pro	Arg	Pro	Ala	Gly	Asp	Ser	Pro	Arg	Lys	Thr	Gln	Trp	Leu	Asn
			100					105					110		
Gln	Val	Glu	Ser	Tyr	lle	ЛÌа	Glu	Gln	Arg	Arg	Gly	Asp	Arg	Met	Arg
		115					120					125			
Pro	Gln	Ala	Pro	Gly	Arg	Gly	Trp	His	Gly	Glu	Glu	Glu	Val	Val	Ala
	130					135					140				
Ala	Ala	Gly	Gln	Glu	Gly	Gln	Val	Glu	Gly	Glu	Glu	Glu	Gly	Glu	Glu
145					150					155					160
Glu	Glu	Glu	Glu	Glu	Asp	Met	Ser	Glu	Val	Phe	Glu	Tyr	Va]	Pro	Va]
				165					170					175	
Phe	Asp	Pro	Val	Val	Asn	Trp	Asp	Gln	Thr	Phe	Ser	Ala	Arg	Asn	Leu
			180					185					190		
Asp	Phe	Gln	Ala	Leu	Arg	Thr	Asp	Trp	lle	Asp	Leu	Ser	Cys	Asn	Thr
		195					200					205			
Ser	Gly	Asn	Leu	Leu	Leu	Pro	Glu	Gln	Glu	Ala	Leu	Glu	Val	Thr	Arg
	210					215					220				
	Phe	Leu	Lys	Lys		Asn	Gln	Arg	Ser		Gly	Arg	Tyr	Gln	
225					230					235					240
Gln	Arg	lle	Val		Val	Glu	Lys	Arg		Asp	Gln	Leu	Arg		Gly
				245					250					255	
Arg	Tyr	Leu		Glu	Leu	Glu	Leu		Glu	GIn	Gly	GIn		Val	Val
		~	260	m	., .		.,	265	0.1	Tr.	61	0.1	270		rs
Arg	Leu		Glu	lyr	val	Ser		Arg	GIY	irp	GIn		116	Asp	Pro
41.	C1	275	C1	C1	V - 1	C1	280	Δ	Λ	1	C1	285	Lan	V.s.1	Т
Ala		61y	GIU	GIU	vai	Glu	мта	Arg	ASN	Leu		61 y	Leu	v 21 1	rrp
Λ	290 D	11.	Λ	A 20.00	A	295	C1.	V = 1	1	Aan	300	A 12.00	Alo	Cln	C1
-	Pro	HIS	ASII	Arg	310	Arg	GIN	vai	Leu	315	Ш	W1.8	Ата	GIII	320
305 Pro	Lva	Lou	Cvc	Trn		Gln	Clu	Dho	Sor		Sor	Hic	Ara	Δla	
110	LyS	Leu	Cys	325	110	0111	Gry	THE	330	пр	361	1115	лιξ	335	, CI I
Val	Hic	Pho	Val		Pro	Val	Lvc	Aan		Δla	Ara	Trn	Val		Gln
vai	1115	THE	340	vai	110	, (11	Lyo	345	0111	ша	Mg	пþ	350	0111	0111
Phe	He	lve		Met	Glu	Asn] eu		Gln	Val	Thr	Glv		Pro	llic
1 116	110	355	цор	in C C	314		360	. 110	0.111	, a 1	1111	365	,5		

Phe Asn Ile Val Ile Thr Asp Tyr Ser Ser Glu Asp Met Asp Val Glu Met Ala Leu Lys Arg Ser Lys Leu Arg Ser Tyr Gln Tyr Val Lys Leu Ser Gly Asn Phe Glu Arg Ser Ala Gly Leu Gln Ala Gly 11e Asp Leu Val Lys Asp Pro His Ser lle Ile Phe Leu Cys Asp Leu His lle His Phe Pro Ala Gly Val Ile Asp Ala Ile Arg Lys His Cys Val Glu Gly Lys Met Ala Phe Ala Pro Met Val Met Arg Leu His Cys Gly Ala Thr Pro Gln Trp Pro Glu Gly Tyr Trp Glu Val Asn Gly Phe Gly Leu Leu Gly Ile Tyr Lys Ser Asp Leu Asp Arg Ile Gly Gly Met Asn Thr Lys Glu Phe Arg Asp Arg Trp Gly Gly Glu Asp Trp Glu Leu Leu Asp Arg lle Leu Gln Ala Gly Leu Asp Val Glu Arg Leu Ser Leu Arg Asn Phe Phe His His Phe His Ser Lys Arg Gly Met Trp Ser Arg Arg Gln Met Lys Thr Leu <210> 4781 <211> 443 <212> PRT

<400> 4781

<213> Homo sapiens

Met Ser Gln Arg Gln Tyr Pro Glu Cys Tyr Leu Ala Pro Asn Gly Cys

1 5 10 15

Leu Val Ser Asn Cys Gly Val Asn Lys Met Ser Asn Glu Glu Leu Val
20 25 30

Gly	Gln	Asn	His	Gly	Met	Glu	G1 y	Glu	Ala	Cys	Thr		Gly	Asp	Val
		35					40					45			
Thr	Phe 50	Ser	Asp	Val	Ala	11e 55	Asp	Phe	Ser	His	G1u 60	Glu	Trp	Ala	Cys
Leu	Asp	Ser	Ala	Gln	Arg	Asp	Leu	Tyr	Lys	Asp	Val	Met	Val	G1n	Asn
65					70					75					80
Tyr	Glu	Asn	Leu	Val	Ser	Val	Gly	Leu	Ser	Val	Thr	Lys	Pro	Tyr	Val
				85					90					95	
He	Met	Leu	Leu	${\tt Glu}$	Asp	Gly	Lys	Glu	Pro	Trp	Met	Met	Glu	Lys	Lys
			100					105					110		
Leu	Ser	Lys	Asp	Trp	Glu	Ser	Arg	Trp	Glu	Asn	Lys	Glu	Leu	Ser	Thr
		115					120					125			
Lys	Lys	Asp	lle	Tyr	Asp	Glu	Asp	Ser	Pro	Gln	Pro	Val	Thr	Met	Glu
	130					135					140				
Lys	Val	Val	Lys	Gln	Ser	Tyr	Glu	Phe	Ser	Asn	Ser	Asn	Lys	Asn	Leu
145					150					155					160
Glu	Tyr	Thr	Glu	Cys	Asp	Thr	Phe	Arg	Ser	Thr	Phe	His	Ser	Lys	Ser
				165					170					175	
Thr	Leu	Ser	Glu	Pro	Gln	Asn	Asn	Ser	Ala	Glu	Gly	Asn	Ser	His	Lys
			180					185					190		
Tyr	Asp	lle	Leu	Lys	Lys	Asn	Leu	Ser	Lys	Lys	Ser		He	Lys	Ser
		195					200					205			
Glu	Arg	lle	Asn	Gly	Gly	Lys	Lys	Leu	Leu	Asn		Asn	Lys	Ser	Gly
	210					215					220				
	Ala	Phe	Asn	Gln		Lys	Ser	Leu	Thr		Pro	Gln	Thr	Cys	
225					230					235			20-		240
Arg	Glu	Lys	He	-	Thr	Cys	Ser	Glu	-	G1 y	Lys	Ala	Phe		Lys
				245					250					255	
Gln	Ser	He	Leu	Asn	Arg	His	Trp	Arg	He	His	Thr	Gly	Glu	Lys	Pro
			260					265					270		
Tyr	Glu	Cys	Arg	Glu	Cys	G1 y	Lys	Thr	Phe	Ser	His	Gly	Ser	Ser	Leu
		275					280					285			
Thr	Arg	His	Gln	Ile	Ser	His	Ser	Gly	Glu	Lys	Pro	Tyr	Lys	Cys	Пе
	290					295					300				
G1 ₁₁	Cve	Glv	lve	Ala	Pho	Sor	Hie	Glv	Sor	Sor	Leu	Thr	Asn	His	G1n

305	31	0	315	320
			r Glu Cys Met Asn	
Sei III IIIS		u Lys 110 ly		
	325		330	335
Ser Phe Ser I	Arg Val Se	r Leu Leu Il	le Gln His Leu Arg	lle His Thr
;	340	34	15	350
Gln Glu Lys	Arg Tyr Gl	u Cys Arg Il	le Cys Gly Lys Ala	Phe Ile His
355		360	365	
Ser Ser Ser l	Leu Ile Hi	s His Gln Ly	ys Ser His Thr Gly	Glu Lys Pro
370		375	380	
Tyr Glu Cys	Arg Glu Cy	s Gly Lys Al	la Phe Cys Cys Ser	Ser His Leu
385	39	90	395	400
Thr Gln His	Gln Arg Il	e His Ser Me	et Lys Lys Lys Tyr	Glu Cys Asn
	405		410	415
Lys Cys Leu	Lys Val Ph	ne Ser Ser Ph	ne Ser Phe Leu Val	Gln His Gln
	420	42	25	430
Ser lle His	Thr Glu Gl	u Lys Pro Ph	ne Glu Val	
435		440		

<211> 649

<212> PRT

<213> Homo sapiens

<400> 4782

65

Met Met Trp Gln Cys His Leu Ser Ala Gln Asp Tyr Arg Tyr Pro 1 15 1 16 1 17 1 18 1 19 1

75

80

lle	Gly	lle	Leu		Ser	Gly	His	Arg		Arg	lle	Leu	Gln		Ile
				85					90					95	
Gln	Leu	Leu	Pro	Lys	Met	Arg	Pro	He	G1 y	His	Asp	Gly	Tyr	His	Pro
			100					105					110		
Thr	Ser	Val	Ala	Glu	Trp	Leu	Asp	Ser	He	Glu	Leu	Gly	Asp	Tyr	Thr
		115					120					125			
Lys	Ala	Phe	Leu	lle	Asn	Gly	Tyr	Thr	Ser	Met	Asp	Leu	Leu	Lys	Lys
	130					135					140				
He	Trp	Glu	Val	Glu	Leu	Ile	Asn	Glu	Trp	Asn	Ile	Thr	Lys	Leu	Ser
145					150					155					160
He	Glu	Tyr	Asp	Ser	Glu	Pro	Phe	Gly	Lys	Glu	Arg	Asp	Ala	Ala	Ile
				165					170					175	
Lys	Lys	Leu	Ala	Thr	Glu	Ala	Gly	Val	Glu	Val	lle	Val	Arg	lle	Ser
			180					185					190		
His	Thr	Leu	Tyr	Asp	Leu	Asp	Lys	Ile	Ile	Glu	Leu	Asn	Gly	Gly	Gln
		195					200			•		205			
Pro	Pro	Leu	Thr	Tyr	Lys	Arg	Phe	Gln	Thr	Leu	He	Ser	Lys	Met	Glu
	210					215					220				
Pro	Leu	Glu	Ile	Pro	Val	Glu	Thr	He	Thr	Ser	Glu	Val	Ile	Glu	Lys
225					230					235					240
Cys	Thr	Thr	Pro	Leu	Ser	Asp	Asp	His	Asp	Glu	Lys	Tyr	Gly	Val	Pro
				245					250					255	
Ser	Leu	Glu	Glu	Leu	Gly	Phe	Asp	Thr	Asp	Gly	Leu	Ser	Ser	Ala	Val
			260					265					270		
Trp	Pro	Gly	Gly	Glu	Thr	Glu	Ala	Leu	Thr	۸rg	Leu	Glu	Arg	His	Leu
		275					280					285			
Glu	Arg	Lys	Ala	Trp	Val	Ala	Asn	Phe	Glu	Arg	Pro	Arg	Met	Asn	Ala
	290					295					300				
Asn	Ser	Leu	Leu	Ala	Ser	Pro	Thr	Gly	Leu	Ser	Pro	Tyr	Leu	Arg	Phe
305					310					315					320
Gly	Cys	Leu	Ser	Cys	Arg	Leu	Phe	Tyr	Phe	Lys	Leu	Thr	Asp	Leu	Tyr
				325					330					335	
Lys	Lys	Val	Lys	Lys	Asn	Ser	Ser	Pro	Pro	Leu	Ser	Leu	Tyr	G1 y	Gln
			340					345					350		
Leu	Leu	Trp	Arg	Glu	Phe	Phe	Tyr	Thr	Ala	Ala	Thr	Asn	Asn	Pro	Arg
		355					360					365			

Phe	Asp	Lys	Met	Glu	Gly	Asn	Pro	He	Cys	Val	Gln	lle	Pro	Trp	Asp
	370					375					380				
Lys	Asn	Pro	Glu	Ala	Leu	Ala	Lys	Trp	Ala	Glu	Gly	Arg	Thr	Gly	Phe
385					390					395					400
Pro	Trp	11e	Лsp	Ala	lle	Met	Thr	Gln	Leu	Arg	Gln	Gl u	Gly	Trp	He
				405					410					415	
His	His	Leu	Ala	Arg	His	Ala	Val	Ala	Cys	Phe	Leu	Thr	Arg	Gly	Asp
			420					425					430		
Leu	Trp	Ile	Ser	Trp	Glu	Glu	Gly	Met	Lys	Val	Phe	Glu	Glu	Leu	Leu
		435					440					445			
Leu	Asp	Ala	Asp	Trp	Ser	He	Asn	Ala	Gly	Ser	Trp	Met	Trp	Leu	Ser
	450					455					460				
Cys	Ser	Ser	Phe	Phe	Gln	Gln	Phe	Phe	His	Cys	Tyr	Cys	Pro	Val	G1 y
465					470					475					480
Phe	Gly	Arg	Arg	Thr	Asp	Pro	Asn	Gly	Asp	Tyr	He	Arg	Arg	Tyr	Leu
				485					490					495	
Pro	Val	Leu	Arg	Gly	Phe	Pro	Ala	Lys	Tyr	11e	Tyr	Asp	Pro	Trp	Asn
			500					505					510		
Ala	Pro	Glu	Gly	Ile	Gln	Lys	Val	Ala	Lys	Cys	Leu	He	G1 y	Val	Asn
		515					520					525			
Tyr	Pro	Lys	Pro	Met	Val	Asn	His	Ala	Glu	Ala	Ser	Arg	Leu	Asn	He
	530					535					540				
Glu	Arg	Met	Lys	Gln	He	Tyr	Gln	G1n	Leu	Ser	Arg	Tyr	Arg	Gly	Leu
545					550					555					560
Gly	Leu	Leu	Ala	Ser	Val	Pro	Ser	Asn	Pro	Asn	Gly	Asn	Gly	Gly	Phe
				565					570					575	
Met	Gly	Tyr	Ser	Ala	Glu	Asn	lle			Cys	Ser	Ser			Ser
			580					585					590		
Cys	Ser		Gly	Ser	Gly	He		His	Tyr	Ala	His		Asp	Ser	Gln
		595					600					605			
Gln		His	Leu	Leu	Lys		Gly	Arg	Ser	Ser			Thr	Gly	Leu
	610					615					620				
	Gly	Gly	Lys	Arg	Pro	Ser	Gln	Glu	Glu			Gln	Ser	lle	
625					630	_				635					640
Pro	Lys	Val	Gln		Gln	Ser	Thr	Asn							
				645											

```
<211> 337
<212> PRT
<213> Homo sapiens
<400> 4783
Met Pro Gly Leu Pro Cys Pro Ala Leu Pro Cys Pro Ala Pro Pro Pro
 1
                  5
                                     10
                                                          15
Ala Pro Ser Cys Leu Ile Ala Met Ala Leu Lys Asn Ser Lys Thr Gly
                                 25
Ser Leu Pro Val Ser Glu 11e Tyr Ser Phe Met Lys Glu His Phe Pro
                             40
                                                  45
Tyr Phe Lys Thr Ala Pro Asp Gly Trp Lys Asn Ser Val Arg His Asn
     50
                         55
                                             60
Leu Ser Leu Asn Lys Cys Phe Glu Lys Val Glu Asn Lys Met Ser Gly
                     70
                                         75
Ser Ser Arg Lys Gly Cys Leu Trp Ala Leu Asn Leu Ala Arg lle Asp
                 85
                                     90
Lys Met Glu Glu Glu Met His Lys Trp Lys Arg Lys Asp Leu Ala Ala
                                105
lle His Arg Ser Met Ala Asn Pro Glu Glu Leu Asp Lys Leu lle Ser
        115
                            120
                                                 125
Asp Arg Pro Glu Ser Cys Arg Arg Pro Gly Lys Pro Gly Glu Pro Glu
    130
                        135
Ala Pro Val Leu Thr His Ala Thr Thr Val Ala Val Ala His Gly Cys
                    150
                                        155
Leu Ala Val Ser Gln Leu Pro Pro Gln Pro Leu Met Thr Leu Ser Leu
                                    170
Gln Ser Val Pro Leu His His Gln Val Gln Pro Gln Ala His Leu Ala
                                185
Pro Asp Ser Pro Ala Pro Ala Gln Thr Pro Pro Leu His Ala Leu Pro
        195
                            200
                                                 205
Asp Leu Ser Pro Ser Pro Leu Pro His Pro Ala Met Gly Arg Ala Pro
   210
                        215
                                             220
```

Val Asp Phe Ile Asn Ile Ser Thr Asp Met Asn Thr Glu Val Asp Ala Leu Asp Pro Ser 11e Met Asp Phe Ala Leu Gln Gly Asn Leu Trp Glu Glu Met Lys Asp Glu Gly Phe Ser Leu Asp Thr Leu Gly Ala Phe Ala Asp Ser Pro Leu Gly Cys Asp Leu Gly Ala Ser Gly Leu Thr Pro Ala Ser Gly Gly Ser Asp Gln Ser Phe Pro Asp Leu Gln Val Thr Gly Leu Tyr Thr Ala Tyr Ser Thr Pro Asp Ser Val Ala Ala Ser Gly Thr Ser Ser Ser Ser Gln Tyr Leu Gly Ala Gln Gly Asn Lys Pro Ile Ala Leu Leu

<210> 4784

<211> 594

<212> PRT

<213> Homo sapiens

<400> 4784

Met Glu Cys Tyr Tyr Ile Val Ile Ser Ser Thr His Leu Ser Asn Gly His Phe Arg Asn 11e Lys Gly Val Phe Arg Gly Pro Leu Ser Lys Asn Gly Asn Lys Thr Leu Asp Tyr Ala Glu Lys Glu Asn Thr Ile Ala Lys Ala Leu Glu Asp Leu Lys Ala Asn Phe Tyr Cys Glu Leu Cys Asp Lys Gln Tyr Tyr Lys His Gln Glu Phe Asp Asn His Ile Asn Ser Tyr Asp His Ala His Lys Gln Arg Leu Lys Glu Leu Lys Gln Arg Glu Phe Ala

Arg	Asn	Val	Ala	Ser	Lys	Ser	Arg	Lys	Asp	Glu	Arg	Lys	Gln	Glu	Lys
			100					105					110		
Ala	Leu	Gln	Arg	Leu	His	Lys	Leu	Ala	Glu	Leu	Arg	Lys	Glu	Thr	Val
		115					120					125			
Cys	Λla	Pro	Gly	Ser	Gly	Pro	Met	Phe	Lys	Ser	Thr	Thr	Val	Thr	Val
	130					135					140				
Arg	Glu	Asn	Cys	Asn	Glu	He	Ser	Gln	Arg	Val	Val	Val	Asp	Ser	Val
145					150					155					160
Asn	Asn	Gln	Gln	Asp	Phe	Lys	Tyr	Thr	Leu	Ile	His	Ser	Glu	Glu	Asn
				165					170					175	
Thr	Lys	Asp	Ala	Thr	Thr	Val	Ala		Asp	Pro	Glu	Ser		Asn	Asn
			180					185					190		
Tyr	Thr		Lys	Asn	Asn	Gln		G1 y	Asp	Gln	Ala		Gly	He	His
		195				_	200		5 1			205			
Arg		Lys	He	Gly	Phe		Phe	Ala	Phe	Pro		Lys	Ala	Ser	Vai
,	210	C.I	C	C	A 7	215	4.3	DI	C	C1	220 T	C	۸	۸	۸1
	Leu	Glu	Ser	Ser		Ala	Ala	Pne	ser		ıyr	ser	ASP	Asp	
225	V = 1	C1	Lua	C1	230	Con	Ana	Lva	Son	235	Dho	Vol	Dro	Sor	240
261.	vai	GIY	Lys	245	rne	Sei	MI B	LyS	250	MIG	THE	vai	110	Ser 255	піа
Cve	Hic	Lau	Gln		Ser	Ser	Pro	Thr		Val	Leu	Len	Ser	Ser	Glu
Cys	1113	Lea	260	Leu	561	DCI	110	265	пор	, (1)	Leu	ДСС	270	561	oru
GIn	lvs	Thr		Ser	Phe	His	Pro		Glu	Ala	Met	Cvs		Asp	Lvs
014	13,0	275			•		280					285	6		, -
Glu	Thr		G1n	Thr	Gln	Glu		Lys	Glu	Val	Ser		Glu	Lys	Лsр
	290					295					300				
Ala		Leu	Leu	Pro	Ser	Phe	Cys	Lys	Phe	Gln	Leu	Gln	Leu	Ser	Ser
305					310					315					320
Asp	Ala	Asp	Asn	Cys	Gln	Asn	Ser	Val	Pro	Leu	Ala	Asp	Gln	He	Pro
				325					330					335	
Leu	Glu	Ser	Val	Val	He	Asn	Glu	Asp	He	Pro	Val	Ser	Gly	Asn	Ser
			340					345					350		
Phe	Glu	Leu	Leu	Gly	Asn	Lys	Ser	Thr	Va]	Leu	Asp	Met	Ser	Asn	Asp
		355					360					365			
Cys	He	Ser	Val	Gln	Ala	Thr	Thr	Glu	Glu	Asn	Val	Lys	His	Asn	Glu
	370					375					380				

Ala	Ser	Thr	Thr	Glu	Val	Glu	Asn	Lys	Asn	Gly	Pro	Glu	Thr	Leu	Ala
385					390					395					400
Pro	Ser	Asn	Thr	Glu	Glu	Va]	Asn	He	Thr	He	His	Lys	Lys	Thr	Asn
				405					410					415	
Phe	Cys	Lys	Arg	Gln	Cys	Glu	Pro	Phe	Val	Pro	Val	Leu	Asn	Lys	His
			420					425					430		
Arg	Ser	Thr	Val	Leu	Gln	Trp	Pro	Ser	Glu	Met	Leu	Val	Tyr	Thr	Thr
		435					440					445			
Thr	Lys	Pro	Ser	lle	Ser	Tyr	Ser	Cys	Asn	Pro	Leu	Cys	Phe	Asp	Phe
	450					455					460				
Lys	Ser	Thr	Lys	Val	Asn	Asn	Asn	Leu	Asp	Lys	Asn	Lys	Pro	Asp	Leu
465					470					475					480
Lys	Asp	Leu	Cys	Ser	Gln	Gln	Lys	Gln	Glu	Asp	lle	Cys	Met	Gly	Pro
				485					490					495	
Leu	Ser	Asp	Tyr	Lys	Asp	Val	Ser	Thr	Glu	Gly	Leu	Thr	Asp	Tyr	Glu
			500					505					510		
11e	Gly	Ser	Ser	Lys	Asn	Lys	Cys	Ser	Gln	Val	Thr	Pro	Leu	Leu	Ala
		515					520					525			
Asp	Asp	lle	Leu	Ser	Ser	Ser	Cys	Asp	Ser	Gly	Lys	Asn	Lys	Asn	Thr
	530					535					540				
Gly	Gln	Arg	Tyr	Lys	Asn	He	Ser	Cys	Lys	lle	Arg	Glu	Thr	Glu	Lys
545					550					555					560
Tyr	Asn	Phe	Thr	Lys	Ser	Gln	He	Lys	Gln	Asp	Thr	Leu	Asp	Glu	Lys
				565					570					575	
Tyr	Asn	Lys	He	Arg	Leu	Lys	Glu	Thr	His	Glu	Tyr	Trp	Phe	His	Lys
			580					585					590		
Ser	Arg														

<211> 118

<212> PRT

<213> Homo sapiens

<400> 4785

Met	11e	Thr	Phe	Tyr	Phe	Leu	Phe	Leu	Phe	Ser	Phe	Leu	Ser	Phe	Phe
1				5					10					15	
Phe	Phe	Phe	Phe	Phe	Glu	Thr	Glu	Phe	Cys	Ser	Ser	Cys	Pro	Gly	Trp
			20					25					30		
Ser	Asn	Gly	Thr	He	Leu	Ala	His	Arg	Asn	Leu	His	Leu	Pro	Gly	Ser
		35					40					45			
Ser	Asp	Ser	Pro	Pro	Ser	Ala	Phe	Pro	Ser	Ser	Trp	Asp	Tyr	Arg	His
	50					55					60				
Ala	Pro	Pro	Ser	Pro	Ala	Asn	Phe	Val	Phe	Leu	Val	Asp	Met	Gly	Leu
65					70					75					80
Leu	His	Val	Gly	Gln	Ala	Gly	Leu	Glu	Leu	Pro	Thr	Ser	Gly	Asp	Pro
				85					90					95	
Pro	Thr	Ser	Ala	Ser	Gln	Ser	Ala	Gly	lle	Thr	Gly	Val	Ser	His	Cys
			100					105					110		
Ala	Arg	Pro	Leu	Phe	Ser										
		115													
<21	0> 47	786													
<21	1> 49	91													
<21	2> PI	RT													
<21	3> Ho	omo s	sapie	ens											
<40	0> 47	786													
Met	Ala	Tyr	Leu	Tyr	Ser	Ser	Asp	Ala	Phe	Leu	Glu	Gly	Tyr	Val	Gln
1				5					10					15	
Gln	Phe	Leu	Tyr	Thr	Phe	Arg	Tyr	Phe	Cys	Thr	Pro	His	Asp	Phe	Leu
			20					25					30		
His	Phe	Leu	Leu	Asp	Arg	He	Asn	Ser	Thr	Leu	Thr	Arg	Ala	His	Gln
		35					40					45			
Asp		Thr	Ser	Thr	Phe	Thr	Lys	Пe	Tyr	Arg	Arg	Ser	Leu	Cys	Val
	Pro	1 1 1 1 1	CC.												
	Pro 50	1111				55					60				
Leu					Glu		Cys	Tyr	Ala	Val		Phe	Pro	Arg	Asn

Ser Gly Leu Leu Gly Lys Leu Glu Asp Phe IIe Ser Ser Lys IIe Leu

Pro	Leu	Asp	Gly	Ser	Ala	Lys	His	Leu	Leu	Gly	Leu	Leu	Glu	Val	Gly
			100					105					110		
Met	Asp	Arg	۸rg	Ala	Glu	Gly	Asn	Pro	Arg	Gly	Thr	Asp	Leu	Glu	Asn
		115					120					125			
Pro	Arg	Glu	Ala	Glu	Glu	Asp	Ala	Arg	Pro	Phe	Asn	Ala	Leu	Cys	Lys
	130					135					140				
Arg	Leu	Ser	Glu	Asp	Gly	He	Ser	Arg	Lys	Ser	Phe	Pro	Trp	Arg	Leu
145					150					155					160
Pro	Arg	Gly	Asn	Gly	Leu	Val	Leu	Pro	Pro	His	Lys	Glu	Arg	Pro	Tyr
				165					170					175	
Thr	He	Ala	Ala	Ala	Leu	Pro	Lys	Pro	Cys	Phe	Leu	Glu	Asp	Phe	Tyr
			180					185					190		
G1 y	Pro	Cys	Ala	Lys	Thr	Ser	Glu	Lys	Gly	Pro	Tyr	Phe	Leu	Thr	Glu
		195					200					205			
Tyr	Ser	Thr	His	Gln	Leu	Phe	Ser	Gln	Leu	Thr	Leu	Leu	Gln	Gln	Glu
	210					215					220				
Leu	Phe	Gln	Lys	Cys	His	Pro	Val	His	Phe	Leu	Asn	Ser	Arg	Ala	Leu
225					230					235					240
Gly	Val	Met	Asp	Lys	Ser	Thr	Ala	He	Pro	Lys	Ala	Ser	Ser	Ser	Glu
				245					250					255	
Ser	Leu	Ser	Ala	Lys	Thr	Cys	Ser	Leu	Phe	Leu	Pro	Asn	Tyr	Val	G1n
			260					265					270		
Asp	Lys	Tyr	Leu	Leu	Gln	Leu	Leu	Arg	Asn	Ala	Asp	Asp	Val	Ser	Thr
		275					280					285			
Trp	Val	Ala	Ala	Glu	He	Val	Thr	Ser	His	Thr	Ser	Lys	Leu	Gln	Val
	290					295					300				
Asn	Leu	Leu	Ser	Lys	Phe	Leu	Leu	He	Ala	Lys	Ser	Cys	Tyr	Glu	Gln
305					310					315					320
Arg	Asn	Phe	Ala	Thr	Ala	Met	GIn	He		Ser	Gly	Leu	Glu		Leu
				325					330					335	
Ala	Val	Arg	Gln	Ser	Pro	Ala	Trp	Arg	He	Leu	Pro	Ala	Lys	He	Ala
			340					345					350		
Glu	Val	Met	Glu	Glu	Leu	Lys	Ala	Val	G1u	Va]	Phe	Leu	Lys	Ser	Asp
		355					360					365			
Ser	Leu	Cys	Leu	Met	Glu	G1 y	Arg	Arg	Phe	Arg	Ala	Gln	Pro	Thr	Leu
	370					375					380				

Pro Ser Ala His Leu Leu Ala Met His lle Gln Gln Leu Glu Thr Gly Gly Phe Thr Met Thr Asn Gly Ala His Arg Trp Ser Lys Leu Arg Asn Ile Ala Lys Val Val Ser Gln Val His Ala Phe Gln Glu Asn Pro Tyr Thr Phe Ser Pro Asp Pro Lys Leu Gln Ser Tyr Leu Lys Gln Arg Ile Ala Arg Phe Ser Gly Ala Asp Ile Ser Thr Leu Ala Ala Asp Ser Arg Ala Asn Phe His Gln Val Ser Ser Glu Lys His Ser Arg Lys Ile Gln Asp Lys Leu Arg Arg Met Lys Ala Thr Phe Gln

<210> 4787

<211> 415

<212> PRT

<213> Homo sapiens

<400> 4787

Met Val Ala Tyr Ser Val Gln Val Leu Ala Val Phe Ile Ser Cys Ala lle Leu Thr Leu Ala Met Lys lle Ala Trp Ile Phe Gly Leu Asn Ser Val Gln Asn Ile Thr Ala Asn Leu Ser Val Asp Gly Ser Thr Ser Gly Asn Pro Ile Gln Lys Trp Lys Arg Lys lle Asp Ala Asn Cys Thr Ala Arg Leu Arg Thr Leu Asn Phe Phe Phe Ala Met Ser Gly Lys Val Lys Asp Gly Thr Pro Cys Ser Pro Asn Lys Asn Asp Val Cys lle Asp Gly

Val Cys Glu Leu Val Gly Cys Asp His Glu Leu Gly Ser Lys Ala Val

Ser	Asp	Ala	Cys	G1 y	Val	Cys	Lys	Gly	Asp	Asn	Ser	Thr	Cys	Lys	Phe
		115					120					125			
Tyr	Lys	Gly	Leu	Tyr	Leu	Asn	Gln	His	Lys	Ala	Asn	Glu	Tyr	Tyr	Pro
	130					135					140				
Val	Val	Leu	He	Pro	Ala	G1 y	Ala	Arg	Ser	Пе	Glu	11e	Gln	Glu	Leu
145					150					155					160
Gln	Val	Ser	Ser	Ser	Tyr	Leu	Ala	Val	Arg	Ser	Leu	Ser	Gln	Lys	His
				165					170					175	
Tyr	Leu	Thr	Gly	Gly	Trp	Ser	He	Asp	Trp	Pro	Gly	Glu	Phe	Pro	Phe
			180					185					190		
Ala	Gly	Thr	Thr	Phe	Glu	Tyr	Gln	Arg	Ser	Phe	Asn	Arg	Pro	Glu	Arg
		195		•			200					205			
Leu	Tyr	Ala	Pro	Gly	Pro	Thr	Asn	Glu	Thr	Leu	Val	Phe	Glu	Пе	Leu
	210					215					220				
Met	Gln	Gly	Lys	Asn	Pro	Gly	lle	Ala	Trp	Lys	Tyr	Ala	Leu	Pro	Lys
225					230					235					240
Val	Met	Asn	Gly	Thr	Pro	Pro	Ala	Thr	Lys	Arg	Pro	Ala	Tyr	Thr	Trp
				245					250					255	
Ser	Ile	Val	Gln	Ser	Glu	Cys	Ser	Val	Ser	Cys	Gly	Gly	Gly	Tyr	lle
			260					265					270		
Asn	Val	Lys	Ala	He	Cys	Leu	Arg	Asp	Gln	Asn	Thr	Gln	Val	Asn	Ser
		275					280					285			
Ser	Phe	Cys	Ser	Ala	Lys	Thr	Lys	Pro	Val	Thr	G]u	Pro	Lys	lle	Cys
	290					295					300				
Asn	Arg	Arg	Ala	Cys	Pro	Ala	His	Pro	Val	Tyr	Asn	Met	Val	Ala	Gly
305					310					315					320
Trp	Tyr	Ser	Leu	Pro	Trp	G1n	Gln	Cys	Thr	Val	Thr	Cys	Gly	Gly	Gly
				325					330					335	
Val	Gln	Thr	Arg	Ser	Val	His	Cys	Val	Gln	Gln	Gly	Arg	Pro	Ser	Ser
			340					345					350		
Ser	Cys	Leu	Leu	His	G1n	Lys	Pro	Pro	Val	Leu	Arg	Ala	Cys	Asn	Thr
		355					360					365			
Asn	Phe	Cys	Pro	Ala	Pro	Glu	Lys	Arg	Glu	Asp	Pro	Ser	Cys	Val	Asp
	370					375					380				
Phe	Phe	Asn	Trp	Cys	His	Leu	Val	Pro	Gln	His	Gly	Val	Cys	Asn	His
385					390					395					400

Lys Phe Tyr Gly Lys Gln Cys Cys Lys Ser Cys Thr Arg Lys 11e 405 410 415 <210> 4788 <211> 974 <212> PRT <213> Homo sapiens <400> 4788 Met Arg Arg Ala Gly Ile Gly Glu Asp Ser Arg Leu Gly Leu Gln Ala 10 Gln Pro Gly Ala Glu Pro Ser Pro Gly Arg Ala Gly Thr Glu Arg Ser 30 25 Leu Gly Gly Ala Gln Gly Pro Gly Gln Pro Cys Ser Cys Pro Gly Ala 40 Met Ala Ser Ala Val Arg Gly Ser Arg Pro Trp Pro Arg Leu Gly Leu 55 60 Gln Leu Gln Phe Ala Ala Leu Leu Leu Gly Thr Leu Ser Pro Gln Val 70 75 80 65 His Thr Leu Arg Pro Glu Asn Leu Leu Leu Val Ser Thr Leu Asp Gly 85 90 Ser Leu His Ala Leu Ser Lys Gln Thr Gly Asp Leu Lys Trp Thr Leu 105 110 Arg Asp Asp Pro Val lle Glu Gly Pro Met Tyr Val Thr Glu Met Ala 125 115 120 Phe Leu Ser Asp Pro Ala Asp Gly Ser Leu Tyr Ile Leu Gly Thr Gln 135 140

 Lys Gln Gln Gln Gly Leu Met
 Lys Leu Pro Phe
 Phe Thr Ile Pro Glu Leu Val

 145
 150
 155
 160

 His Ala Ser Pro Cys Arg Ser Ser Asp Gly Val Phe Tyr Thr Gly Arg
 165
 170
 175

 Lys Gln Asp Ala Trp Phe Val Val Asp Pro Glu Ser Gly Glu Thr Gln
 185
 190

 Met Thr Leu Thr Thr Glu Gly Pro Ser Thr Pro Arg Leu Tyr Ile Gly
 195

Arg	Thr	Gln	Tyr	Thr	Val	Thr	Met	His	Asp	Pro	Arg	Ala	Pro	Ala	Leu
	210					215					220				
Arg	Trp	Asn	Thr	Thr	Tyr	Arg	Arg	Tyr	Ser	Ala	Pro	Pro	Met	Asp	Gly
225					230					235					240
Ser	Pro	Gly	Lys	Tyr	Met	Ser	His	Leu	Ala	Ser	Cys	Gly	Met	Gly	Leu
				245					250					255	
Leu	Leu	Thr	Val	Asp	Pro	Gly	Ser	Gly	Thr	Val	Leu	Trp	Thr	G1n	Asp
	-		260					265					270		
Leu	Gly	Val	Pro	Val	Met	Gly	Val	Tyr	Thr	Trp	His	Gln	Asp	G1 y	Leu
		275					280					285			
Arg	Gln	Leu	Pro	His	Leu	Thr	Leu	Ala	Arg	Asp	Thr	Leu	His	Phe	Leu
	290					295					300				
Ala	Leu	Arg	Trp	Gly	His	Ile	Arg	Leu	Pro	Ala	Ser	Gly	Pro	Arg	Asp
305					310					315					320
Thr	Ala	Thr	Leu	Phe	Ser	Thr	Leu	Asp	Thr	Gln	Leu	Leu	Met	Thr	Leu
				325					330					335	
Tyr	Val	Gly	Lys	Asp	Glu	Thr	Gly	Phe	Tyr	Val	Ser	Lys	Ala	Leu	Val
			340					345					350		
His	Thr	Gly	Val	Ala	Leu	Val	Pro	Arg	Gly	Leu	Thr	Leu	Ala	Pro	Ala
		355					360					365			
Asp	Gly	Pro	Thr	Thr	Asp	Glu	Val	Thr	Leu	Gln	Val	Ser	Gly	Glu	Arg
	370					375					380				
Glu	Gly	Ser	Pro	Ser	Thr	Ala	Val	Arg	Tyr	Pro	Ser	G1y	Ser	Val	Ala
385					390					395					400
Leu	Pro	Ser	Gln	Trp	Leu	Leu	He	G1 y	His	His	Glu	Leu	Pro	Pro	Val
				405					410					415	
Leu	His	Thr	Thr	Met	Leu	Arg	Val		Pro	Thr	Leu	Gly	Ser	Gly	Thr
			420					425					430		
Ala	Glu	Thr	Arg	Pro	Pro	Glu	Asn	Thr	Gln	Ala	Pro		Phe	Phe	Leu
		435					440					445			
Glu	Leu	Leu	Ser	Leu	Ser	Arg	Glu	Lys	Leu	Trp	Asp	Ser	G] u	Leu	His
	450					455					460				
Pro	Glu	Glu	Lys	Thr	Pro	Asp	Ser	Tyr	Leu		Leu	G]y	Pro	G1n	
465					470					475					480
Leu	Leu	Ala	Ala		Leu	Thr	Ala	Val		Leu	Gly	Gly	Trp		Leu
				485					490					495	

Phe	Val	Met	Arg	Gln	Gln	Gln	Pro	Gln	Val	Val	Glu	Lys	Gln	Gln	Glu
			500					505					510		
Thr	Pro	Leu	Ala	Pro	Λla	Asp	Phe	Ala	His	He	Ser	Gln	Asp	Ala	Gln
		515					520					525			
Ser	Leu	His	Ser	Gly	Ala	Ser	Arg	Arg	Ser	Gln	Lys	Arg	Leu	Gln	Ser
	530					535					540				
Pro	Ser	Lys	Gln	Ala	Gln	Pro	Leu	Asp	Asp	Pro	Glu	Ala	Glu	G1n	Leu
545					550					555					560
Thr	Val	Val	Gly	Lys	He	Ser	Phe	Asn	Pro	Lys	Asp	Val	Leu		Arg
				565					570					575	
Gly	Ala	G1 y	Gly	Thr	Phe	Val	Phe		Gly	Gln	Phe	Glu	Gly	Arg	Ala
			580					585					590		
Val	Ala		Lys	Arg	Leu	Leu		Glu	Cys	Phe	Gly		Val	Arg	Arg
		595					600					605		_	
Glu		Gln	Leu	Leu	Gln		Ser	Asp	Arg	His		Asn	Val	Leu	Arg
_	610					615	_	0.7	D1		620 m	- 1		,	0.1
	Phe	Cys	Thr	Glu		Gly	Pro	Gln	Phe		Tyr	He	Ala	Leu	
625	_				630	0.1	0.1	4 5	17 1	635		D			640
Leu	Cys	Arg	Ala		Leu	GIn	Glu	Tyr	Val	Glu	Asn	Pro	Asp		Asp
	0.1	C1		645	n	C1.	17 1	17 1	650	C1	C1	1	Mad	655	C1.
Arg	61 y	Gly		GIU	Pro	61 u	vai		Leu	GIN	UIN	Leu		ser	Gly
,	4.1	11.2	660	117.	C	1	112 -	665	V - 1	u: _	Λ	A = 10	670	Luc	Dane
Leu	АТА	675	Leu	HIS	ser	Leu	680	116	Val	піѕ	AI g	685	Leu	Lys	FIC
C1	Aan		Lou	110	Thr	C1v		Aan	Ser	Cln.	Cly		G1 _v	Δra	Val
GIY	690		Leu	116	1111	695	110	nsp	361	UIII	700	Leu	Oly	Mg	141
Val			Asn	Pho	Glv		Cvs	lve	Lys	Len		Ala	Glv	Arø	Cvs
705		001	ЛЭР	THE	710		0,5	Lyo	Lyo	715		111 G	u1,		720
		Ser	Leu	His			He	Pro	Gly			Glv	Trp	Met	
001	1 110	001	200	725		9.,			730					735	
Pro	Glu	Leu	Leu			Leu	Pro	Pro	Asp		Pro	Thr	Ser		Va]
			740					745	•				750		
Asp	He	Phe	Ser	Ala	Glv	Cvs	Val		Tyr	Tyr	Val	Leu	Ser	Gly	Gly
•		755			J	-	760		-	-		765			
Ser	His			Gly	Asp	Ser	Leu	Tyr	Arg	Gln	Ala	Asn	lle	Leu	Thi
	770					775					780				

Gly Ala Pro Cys Leu Ala His Leu Glu Glu Glu Val His Asp Lys Val Val Ala Arg Asp Leu Val Gly Ala Met Leu Ser Pro Leu Pro Gln Pro Arg Pro Ser Ala Pro Gln Val Leu Ala His Pro Phe Phe Trp Ser Arg Ala Lys Gln Leu Gln Phe Phe Gln Asp Val Ser Asp Trp Leu Glu Lys Glu Ser Glu Gln Glu Pro Leu Val Arg Ala Leu Glu Ala Gly Gly Cys Ala Val Val Arg Asp Asn Trp His Glu His Ile Ser Met Pro Leu Gln Thr Asp Leu Arg Lys Phe Arg Ser Tyr Lys Gly Thr Ser Val Arg Asp Leu Leu Arg Ala Val Arg Asn Lys Lys His His Tyr Arg Glu Leu Pro Val Glu Val Arg Gln Ala Leu Gly Gln Val Pro Asp Gly Phe Val Gln Tyr Phe Thr Asn Arg Phe Pro Arg Leu Leu His Thr His Arg Ala Met Arg Ser Cys Ala Ser Glu Ser Leu Phe Leu Pro Tyr Tyr Pro Pro Asp Ser Glu Ala Arg Arg Pro Cys Pro Gly Ala Thr Gly Arg

<210> 4789

<211> 794

<212> PRT

<213> Homo sapiens

<400> 4789

Met Glu Gly Ser Leu Ala Gly Ser Leu Ala Ala Pro Asp Arg Pro Gln

1 5 10 15

Gly Pro Glu Arg Leu Pro Gly Pro Ala Pro Arg Glu Asn 11e Glu Gly

			20					25					30		
Gly	Ala	Glu	Ala	Ala	Glu	G1 y	Glu	G1 y	Gly	lle	Phe	Arg	Ser	Thr	Arg
		35					40					45			
Tyr	Leu	Pro	Val	Thr	Lys	Glu	Gly	Pro	Arg	Asp	lle	Leu	Asp	Gly	Arg
	50					55					60				
Gly	Gly	lle	Ser	Val	Ala	Asn	Phe	Asp	Pro	Gly	Thr	Phe	Ser	Leu	Met
65					70					7 5					80
Arg	Cys	Asp	Phe	Cys	Gly	Ala	Gly	Phe	Asp	Thr	Arg	Ala	Gly	Leu	Ser
				85					90					95	
Ser	His	Ala	Arg	Ala	His	Leu	Arg	Asp	Phe	Gly	He	Thr	Asn	Trp	Glu
			100					105					110		
Leu	Thr	Val	Ser	Pro	lle	Asn	He	Leu	Gln	Glu	Leu	Leu	Ala	Thr	Ser
		115					120					125			
Ala	Ala	Glu	Gln	Pro	Pro	Ser	Pro	Leu	Gly	Arg	Glu	Pro	Gly	Gly	Pro
	130					135					140				
Pro	Gly	Ser	Phe	Leu	Thr	Ser	Arg	Arg	Pro	Arg	Leu	Pro	Leu	Thr	Val
145					150					155					160
Pro	Phe	Pro	Pro	Thr	Trp	Ala	Glu	Asp	Pro	G1 y	Pro	Ala	Tyr	Gly	Asp
				165					170					175	
Ala	Ser	Gly	Pro	Glu	Pro	Ala	Arg	Asp	lle	Arg	Cys	Glu	Phe	Cys	Gly
			180					185					190		
Glu	Phe	Phe	Glu	Asn	Arg	Lys	G1 y	Leu	Ser	Ser	His		Arg	Ser	His
		195					200					205			
Leu		Gln	Met	G1 y	Val	Thr	Glu	Trp	Tyr	Val		Gly	Ser	Pro	He
	210					215					220				0.1
		Leu	Arg	Glu				Arg	Arg			Ser	Arg	Pro	Gly
225				10		0.1			Б	235			. 1	,	240
Gly	Pro	Pro	Asn		Pro	Gly	Pro	Ser		Lys	Ala	Leu	Ala		Met
	6.1	C.1	. 1	245	n	C1	C	C	250	C1	A 1 -	A	C	255	C
Met	GIy	Gly		61 y	PTO	61 y	Ser		Leu	GIU	Ala	Arg		PFO	Ser
Λ	Lan	U; a	260	Can	Dno	Lau	416	265	Lvo	Lou	Dro	Dro	270 Pro	Dro	Clv
Asp	Leu		116	Sel.	PTO	Leu		Lys	Lys	Leu	F10	285	110	110	Gly
Son	Pro	275	G1v	Hic	Sor	Pro	280 Thr	۸1.	Sor	Pro	Pro		Thr	Ala	Arg
Set.	290	Leu	оту	11.1.5	261	295	1111	ura	261	110	300	0	1111	ם ניי	111 8
lve		Phe	Pro	Glv	Leu		Ala	Pro	Ser	Leu		lvs	Lvs	Leu	Lvs

305					310					315					320
Pro	Glu	Gln	He	۸rg	Val	Glu	He	Lys	Arg	Glu	Met	Leu	Pro	Gly	Ala
				325					330					335	
Leu	His	Gly	Glu	Leu	His	Pro	Ser	Glu	Gly	Pro	Trp	Gly	Ala	Pro	Arg
			340					345					350		
Glu	Asp	Met	Thr	Pro	Leu	Asn	Leu	Ser	Ser	Arg	Ala	Glu	Pro	Val	Arg
		355					360					365			
Asp	He	Arg	Cys	Glu	Phe	Cys	Gly	Glu	Phe	Phe	Glu	Asn	Arg	Lys	Gly
	370					375					380				
Leu	Ser	Ser	His	Ala	Arg	Ser	His	Leu	Arg	Gln	Met	Gly	Val	Thr	Glu
385					390					395					400
Trp	Ser	Val	Asn	Gly	Ser	Pro	He	Asp	Thr	Leu	Arg	Glu	Île	Leu	Lys
				405					410					415	
Lys	Lys	Ser	Lys	Pro	Cys	Leu	lle	Lys	Lys	Glu	Pro	Pro	Ala	Gly	Λsp
			420					425					430		
Leu	Ala	Pro	Ala	Leu	Ala	Glu	Asp	Gly	Pro	Pro	Thr	Val	Ala	Pro	Gly
		435					440					445			
Pro	Val	Gln	Ser	Pro	Leu	Pro	Leu	Ser	Pro	Leu	Ala	Gly	Arg	Pro	Gly
	450					455					460				
Lys	Pro	Gly	Ala	Gly	Pro	Ala	Gln	Val	Pro	Arg	Glu	Leu	Ser	Leu	Thr
465					470					475					480
Pro	He	Thr	Gly	Ala	Lys	Pro	Ser	Ala	Thr	Gly	Tyr	Leu	Gly	Ser	Val
				485					490					495	
Ala	Ala	Lys	Arg	Pro	Leu	Gln	Glu	Asp	Arg	Leu	Leu	Pro	Ala	Glu	Val
			500					505					510		
Lys	Ala	Lys	Ala	Tyr	lle	Gln	Thr	Glu	Leu	Pro	Phe	Lys	Ala	Lys	Thr
		515					520					525			
Leu	His	Glu	Lys	Thr	Ser	His	Ser	Ser	Thr	Glu	Ala	Cys	Cys	Glu	Leu
	530					535					540				
Cys	Gly	Leu	Tyr	Phe	Glu	Asn	Arg	Lys	Ala	Leu	Ala	Ser	His	Ala	Arg
545					550					555					560
Ala	His	Leu	Arg	Gln	Phe	Gly	Val	Thr	Glu	Trp	Cys	Val	Asn	Gly	Ser
				565					570					575	
Pro	He	Glu	Thr	Leu	Ser	Glu	Trp	11e	Lys	His	Arg	Pro	Gln	Lys	Val
			580					585					590		
Glv	Ala	Tyr	Arg	Ser	Tvr	He	Gln	Glv	Glv	Arg	Pro	Phe	Thr	Lvs	Lys

		595					600					605			
Phe	Arg	Ser	Ala	Gly	His	Gly	Arg	Asp	Ser	Asp	Lys	Arg	Pro	Ser	Leu
	610					615					620				
Gly	Leu	Ala	Pro	Gly	Gly	Leu	Ala	Val	Val	Gly	Arg	Ser	Ala	G1 y	Gly
625					630					635					640
Glu	Pro	Gly	Pro	Glu	Ala	Gly	Arg	Ala	Ala	Asp	Gly	Gly	Glu	Arg	Pro
				645					650					655	
Leu	Ala	Ala	Ser	Pro	Pro	Gly	Thr	Val	Lys	Ala	Glu	Glu	His	Gln	Arg
			660					665					670		
Gln	Asn	Ile	Asn	Lys	Phe	Glu	Arg	Arg	Gln	Ala	Arg	Pro	Pro	Asp	Ala
		675					680					685			
Ser	Ala	Ala	Arg	Gly	Gly	Glu	Asp	Thr	Asn	Asp	Leu	Gln	Gln	Lys	Leu
	690					695					700				
Glu	Glu	Va]	Arg	Gln	Pro	Pro	Pro	Arg	Val	Arg	Pro	Val	Pro	Ser	Leu
705					710					715					720
Val	Pro	Arg	Pro	Pro	Gln	Thr	Ser	Leu	Val	Lys	Phe	Val	Gly	Asn	He
				725					730					735	
Tyr	Thr	Leu	Lys	Cys	Arg	Phe	Cys	Glu	Val	Glu	Phe	Gln	Gly	Pro	Leu
			740					745					750		
Ser	lle	Gln	Glu	Glu	Trp	Val	Arg	His	Leu	Gln	Arg	His	Ile	Leu	Glu
		755					760					765			
Met	Asn	Phe	Ser	Lys	Ala	Asp	Pro	Pro	Pro	Glu	Glu	Ser	Gln	Ala	Pro
	770					775					780				
Gln	Ala	Gln	Thr	Ala	Ala	Ala	Glu	Ala	Pro						
785					790										
)> 4'														
	1> 69														
	2> PI									•					
<21:	3> H	omo	sapi	ens											
)> 4°				_		0-								
Met	Lys	Glu	Phe	Leu	Ser	Thr	Ala	Gln	Gly	Asn	Arg	G]u	Val	Phe	His

Ala Gly Thr Leu Gln Ile His Glu Ser His His Asn Gly Asp Phe Cys

				20					25					30		
7	ſyr	Gln	Asp	Val	Asp	Lys	Asp	Ile	His	Asp	Tyr	Glu	Phe	GIn	Trp	Gln
			35					40					45			
(Glu	Asp	Glu	Arg	Asn	Gly	His	Glu	Ala	Pro	Met	Thr	Lys	He	Lys	Lys
		50					55					60				
I	_eu	Thr	Gly	Ser	Thr	G1u	Arg	Tyr	Asp	G1n	Ser	His	Ala	Arg	Asn	Lys
	65					70					75					80
ŀ	ro	Ile	Lys	Asp	Gln	Leu	Gly	Ser	Ser	Phe	His	Ser	His	Leu	Pro	Glu
					85					90					95	
N	Met	His	Ile	Phe	Gln	Thr	Glu	Glu	Lys	He	Asp	Asn	Gln	Val	Val	Lys
				100					105					110		
9	Ser	He	His	Asp	Ala	Ser	Leu	Val	Ser	Thr	Ala	Gln	Arg	lle	Ser	Cys
			115					120					125			
I	Arg	Pro	Lys	Thr	His	Ile	Ser	Asn	Asn	His	Gly	Asn	Asn	Phe	Trp	.Asn
		130					135					140				
		Ser	Leu	Leu	Thr		Lys	Gln	Glu	Val		Met	Arg	Glu	Lys	
	145					150					155					160
]	Phe	Gln	Cys	Asn		Ser	Gly	Lys	Ala		Asn	Tyr	Ser	Ser		Leu
					165					170					175	
1	Arg	Lys	His		lle	He	His	Leu		Asp	Lys	Tyr	Lys		Asp	Val
	0	0.1		180	D.	ă.	0.1		185		,	4.3	C	190		
(Cys	Gly		Leu	Phe	Asn	Gln		Arg	Asn	Leu	Ala		His	Arg	Arg
	0	11.	195	CI	C1	Λ	D	200	1	C	Λ	C1.	205	C1	1	Tl
(Lys		Inr	61y	61u	Asn	Pro	ıyr	Lys	Cys	ASN		cys	оту	Lys	ınr
1	Dl	210	C1	The	C a sa	S.c.	215	The	Cuc	и; -	A 20 C	220	Lan	ш	The	61
	225	ser	GIN	ınr	ser	Ser 230	Leu	ınr	Cys	піѕ	Arg 235		reu	шѕ	mr	240
		Lve	Pro	Tur	Lve		Glu	Glu	Cvs	Asn			Phe	Hie	Phe	
,	oıu	Lys	110	1 y 1	245	Cys	oru	oru	Cys	250	rys	ma	ine	1113	255	Lyo
,	Ser	110	Len	Glu		Hic	Ser	He	He		Thr	Glu	Glu	Lvs		Tvr
	OG]	116	⊾ëu	260	шв	1113	261	116	265	1113	1111	OIU	JIU	270		. , 1
	Lvs	Cvs	Asn		Cvs	Glv	Lys	Thr		Arg	Gln	Lvs	Ser		Leu	Thr
		J, G	275	u	5,5	~ . J	2,0	280				, 0	285			
	Arg	His		Arg	Leu	His	Thr		Glu	Lvs	Pro	Tvr		Cys	Asn	Glu
	6	290		0			295	1		, -		300	,	, -		
	Cys		Lys	Thr	Phe	Ser	His	Lys	Ser	Ser	Leu		Cys	His	His	Arg

305					310					315					320
Leu	His	Thr	Gly	Glu	Lys	Pro	Tyr	Lys	Cys	Asn	Glu	Cys	Gly	Lys	Thr
				325					330					335	
Phe	Ser	His	Lys	Ser	Ser	Leu	Thr	Cys	His	Arg	Arg	Leu	His	Thr	Gly
			340					345					350		
Glu	Lys	Pro	Tyr	Lys	Cys	Glu	Glu	Cys	Asp	Lys	Ala	Tyr	Ser	Phe	Arg
		355					360					365			
Ser	Asn	Phe	Glu	lle	His	Arg	Lys	He	His	Thr	Glu	Asp	Asn	Ala	Tyr
	370					375					380				
Lys	Cys	Asn	Glu	Cys	Gly	Lys	Thr	Phe	Ser	Arg	Thr	Ser	Ser	Leu	Thr
385					390					395					400
Cys	His	Arg	Arg	Arg	His	Thr	Gly	Glu	Gln	Pro	Tyr	Lys	Cys	Glu	Glu
				405					410					415	
Cys	Asp	Lys	Ala	Phe	Arg	Phe	Lys	Ser	Asn	Leu	Glu	Arg	His	Arg	Arg
			420					425					430		
He	His	Thr	Gly	Glu	Lys	Pro	Tyr	Lys	Cys	Asn	Glu	Cys	Gly	Lys	Thr
		435					440					445			
Phe	Ser	Arg	Lys	Ser	Tyr	Leu	Thr	Cys	His	His	Arg	Leu	His	Thr	Gly
	450					455					460				
Glu	Lys	Ala	Tyr	Lys	Cys	Asn	Glu	Cys	Gly	Lys	Thr	Phe	Ser	Trp	Lys
465					470					475					480
Ser	Ser	Leu	Thr	Cys	His	Arg	Arg	Leu	His	Ser	Gly	Glu	Lys	Pro	Tyr
				485					490					495	
Lys	Cys	Lys	Glu	Cys	Gly	Lys	Thr	Phe	Asn	Gln	Gln	Leu	Thr	Leu	Lys
			500					505					510		
Arg	His	Arg	Arg	Leu	His	Ser	Gly	Glu	Asn	Pro	Tyr	Lys	Cys	Glu	Asp
		515					520					525			
Ser	Asp	Lys	Ala	Tyr	Ser	Phe	Lys	Ser	Asn	Leu	Glu	He	His	Gln	Lys
	530					535					540				
Пе	His	Thr	Glu	Glu	Asn	Pro	Tyr	Lys	Cys	Asn	Glu	Cys	Gly	Lys	Thr
545					550					555					560
Phe	Ser	Arg	Thr	Ser	Ser	Leu	Thr	Cys	His	Arg	Arg	Leu	His	Thr	Gly
				565					570					575	
Glu	Lys	Pro	Tyr	Lys	Cys	Glu	Glu	Cys	Asp	Lys	Ala	Phe	Arg	Val	Lys
			580					585					590		

Ser Asn Leu Glu Gly His Arg Arg Ile His Thr Gly Glu Lys Pro Tyr 600 Lys Cys Asn Glu Cys Gly Lys Thr Phe Ser Arg Lys Ser Tyr Phe lle 615 620 610 Cys His His Arg Leu His Thr Gly Glu Lys Pro Tyr Lys Cys Asn Glu 630 635 Cys Gly Lys Asn Phe Ser Gln Lys Ser Ser Leu lle Cys His His Arg 650 645 Leu His Thr Gly Glu Lys Pro Tyr Lys Cys Asn Glu Cys Gly Lys Thr 665 660 Phe Ser Gln Lys Ser Asn Leu Thr Cys His Arg Arg Leu His Thr Gly 675 680 685 Glu Lys Gln Val 690

<210> 4791

<211> 872

<212> PRT

<213> Homo sapiens

<400> 4791

Met Gly lle Val Trp lle Leu Met Ser Val Leu Pro Trp Trp Gly Arg

1 5 10 15

Cys Ala Asp Leu Ala IIe Ala Ser Thr Gln Leu Val Pro Ser Thr Ala 20 25 30

Ser Ala Arg Met Ala Leu Ser Ser Gln Leu Met Gly Arg Thr Val Trp 35 40 45

Thr Pro Met Ser Ala Ser Ala Leu Gln Glu Pro Ala Tyr Gln Ala Leu 50 55 60

Ala Arg Thr Ser Arg Ala Pro Ser Ala Ala Ser Val Pro Met Ala Ser 65 70 75 80

Arg Cys Arg Val Thr Thr Ala Leu lle Ser Thr Ser Ala Gln Arg Ser 85 90 95

Pro Thr Ser Ala Ser Leu Ala Pro Val Pro Thr Ala Leu Gly Ala Ser 100 105 110

Ser	Λla	Ser	Ala	His	Leu	Ala	Leu	Ser	Ser	Leu	Thr	Met	Gly	Thr	Val
		115					120					125			
Ala	Leu	Thr	His	Gly	Arg	Val	Ser	Ala	Ser	Pro	Val	Leu	Arg	Leu	Gly
	130					135					140				
Ser	Ala	۸rg	Cys	Pro	Lys	Leu	Ser	Thr	Pro	Pro	Arg	Pro	Ala	Ala	Ala
145					150					155					160
Ala	Val	Arg	G1 y	Leu	Gly	Arg	Λla	Gly	Glu	Thr	Pro	Ala	Asn	Cys	Val
				165					170					175	
Ser	Arg	Arg	Ala	Val	Pro	Gly	Pro	Asp	Asp	Ser	Arg	Glu	Asp	Val	Asn
			180					185					190		
Glu	Cys	Ala	Glu	Asn	Pro	Gly	Val	Cys	Thr	Asn	Gly	Val	Cys	Val	Asn
		195					200					205			
Thr	Asp	Gly	Ser	Phe	Arg	Cys	Glu	Cys	Pro	Phe	Gly	Tyr	Ser	Leu	Asp
	210					215					220				
Phe	Thr	Gly	lle	Asn	Cys	Va]	Asp	Thr	Asp	Glu	Cys	Ser	Val	Gly	His
225					230					235					240
Pro	Cys	Gly	Gln	Gly	Thr	Cys	Thr	Asn	Val	lle	Gly	Gly	Phe	Glu	Cys
				245					250					255	
Ala	Cys	Ala	Asp	Gly	Phe	Glu	Pro	Gly	Leu	Met	Met	Thr	Cys	Glu	Asp
			260					265					270		
He	Asp	Glu	Cys	Ser	Leu	Asn	Pro	Leu	Leu	Cys	Ala	Phe	Arg	Cys	His
	•	275					280					285			
Asn	Thr	Glu	Gly	Ser	Tyr	Leu	Cys	Thr	Cys	Pro	Ala	Gly	Tyr	Thr	Leu
	290					295					300				
Arg	Glu	Asp	Gly	Λla	Met	Cys	Arg	Asp	Val	Asp	Glu	Cys	Ala	Asp	Gly
305					310					315					320
Gln	G1n	Asp	Cys		Ala	Arg	Gly	Met		Cys	Lys	Asn	Leu	He	Gly
				325					330					335	
Thr	Phe	Ala	-	Val	Cys	Pro	Pro		Met	Arg	Pro	Leu		Gly	Ser
			340					345					350	_	
Gly	Glu		Cys	Thr	Asp	Asp		Glu	Cys	His	Ala		Pro	Asp	Leu
		355					360				_	365	_		
Cys		Asn	Gly	Arg	Cys		Asn	Thr	Ala	Gly		Phe	Arg	Cys	Asp
	370		<i>-</i> -	D.	۵.	375			m)		380	0.7			
-	Asp	Glu	Gly	Phe		Pro	Ser	Pro	Thr		Thr	Glu	Cys	His	
385					390					395					400

He	Arg	Gln	Gly	Pro	Cys	Phe	Ala	Glu	Val	Leu	Gln	Thr	Met	Cys	Arg
				405					410					415	
Ser	Leu	Ser	Ser	Ser	Ser	Glu	Ala	Val	Thr	Arg	Ala	Glu	Cys	Cys	Cys
			420					425					430		
Gly	Gly	Gly	Arg	Gly	Trp	Gly	Pro	Arg	Cys	Glu	Leu	Cys	Pro	Leu	Pro
		435					440					445			
Gly	Thr	Ser	Ala	Tyr	Arg	Lys	Leu	Cys	Pro	His	Gly	Ser	Gly	Tyr	Thr
	450					455					460				
Ala	Glu	Gly	Arg	Asp	Val	Asp	Glu	Cys	Arg	Met	Leu	Ala	His	Leu	Cys
465					470					475					480
Ala	His	Gly	Glu	Cys	lle	Asn	Ser	Leu	Gly	Ser	Phe	Arg	Cys	His	Cys
				485					490					495	
Gln	Ala	Gly	Tyr	Thr	Pro	Asp	Ala	Thr	Ala	Thr	Thr	Cys	Leu	Asp	Met
			500					505					510		
Asp	Glu	Cys	Ser	Gln	Val	Pro	Lys	Pro	Cys	Thr	Phe	Leu	Cys	Lys	Asn
		515					520					525			
Thr	Lys	Gly	Ser	Phe	Leu	Cys	Ser	Cys	Pro	Arg	Gly	Tyr	Leu	Leu	Glu
	530					535					540				
Glu	Asp	Gly	Arg	Thr	Cys	Lys	Asp	Leu	Asp	Glu	Cys	Thr	Ser	Arg	Gln
545					550					555					560
His	Asn	Cys	Gln	Phe	Leu	Cys	Val	Asn	Thr	Val	Gly	Ala	Phe	Thr	Cys
				565					570					575	
Arg	Cys	Pro	Pro	Gly	Phe	Thr	Gln	His	His	Gln	Ala	Cys	Phe	Asp	Val
			580					585					590		
Asn	Glu	Cys	Asp	Gly	Pro	His	Arg	Cys	Gln	His	Gly	Cys	Gln	Asn	Gln
		595					600					605			
Leu	Gly	Gly	Tyr	Arg	Cys	Ser	Cys	Pro	Gln	Gly		Thr	Gln	His	Ser
	610					615					620				
	Trp	Ala	Gln	Cys		Asp	Glu	Asn	G] u		Ala	Leu	Ser	Pro	
625					630					635					640
Thr	Cys	Gly	Ser		Ser	Cys	Arg	Asn		Leu	G1 y	Gly	Phe		Cys
				645					650					655	
Val	Cys	Pro		Gly	Phe	Asp	Phe		Gln	Ala	Leu	Gly		Cys	Gln
			660					665			_	_	670		.~
Asp	Val		Glu	Cys	Ala	G] y		Arg	Gly	Pro	Cys		Tyr	Ser	Cys
		675					680					685			

Ala Asn Thr Pro Gly Gly Phe Leu Cys Gly Cys Pro Gln Gly Tyr Phe Arg Ala Gly Gln Gly His Cys Val Ser Gly Leu Gly Phe Ser Pro Gly Pro Gln Asp Thr Pro Asp Lys Glu Glu Leu Leu Ser Ser Glu Ala Cys Tyr Glu Cys Lys Ile Asn Gly Leu Ser Pro Arg Asp Arg Pro Arg Arg Ser Ala His Arg Asp His Gln Val Asn Leu Ala Thr Leu Asp Ser Glu Ala Leu Leu Thr Leu Gly Leu Asn Leu Ser His Leu Gly Arg Ala Glu Arg Ile Leu Glu Leu Arg Pro Ala Leu Glu Gly Leu Glu Gly Arg Ile Arg Tyr Val lle Val Arg Gly Asn Glu Gln Gly Phe Phe Arg Met His His Leu Arg Gly Val Ser Ser Leu Gln Leu Gly Arg Arg Arg Pro Gly Pro Gly Thr Tyr Arg Leu Glu Val Val Ser His Met Ala Gly Pro Trp Gly Val Gln Pro Glu Gly Gln Pro Gly Pro Trp Gly Gln Ala Leu Arg Leu Lys Val Gln Leu Gln Leu Leu

<210> 4792

<211> 1015

<212> PRT

<213> Homo sapiens

<400> 4792

Met Asp lle Met Ser Ala Gln Ala Ser Ser Gly Ser Thr Ser Val Leu

1 5 10 15

Arg Trp Val Leu Ser Cys Leu Ala Thr Leu Leu Arg Lys Gln Asp Leu

Glu	Ala	Trp	Gly	Tyr	Pro	Val	Thr	Leu	Gln	Val	Tyr	His	Gly	Leu	Leu
		35					40					45			
Ser	Phe	Thr	Val	His	Pro	Lys	Pro	Lys	Leu	Val	Thr	Ala	Cys	Ala	Met
	50					55					60				
Gln	Ala	Phe	His	Ser	Leu	Phe	His	Ala	Arg	Pro	Gly	Leu	Ser	Thr	Leu
65					70					75					80
Ser	Ala	Glu	Leu	Λsn	Ala	Gln	Ile	He	Thr	Λla	Leu	Tyr	Asp	Tyr	Val
				85					90					95	
Pro	Ser	Glu	Asp	Asp	Leu	Gln	Pro	Leu	Leu	Ala	Trp	Leu	Lys	Val	Met
			100					105					110		
Glu	Lys	Ala	His	Ile	Așn	Leu	Val	Arg	Leu	Gln	Trp	Asp	Leu	Gly	Leu
		115					120					125			
Gly	His	Leu	Pro	Arg	Phe	Phe	Gly	Thr	Ala	Val	Thr	Cys	Leu	Leu	Ser
	130					135					140				
Pro	His	Ser	Gln	Val	Leu	Thr	Ala	Ala	Thr	Gln	Ser	Leu	Lys	Glu	Jle
145					150					155					160
Leu	Lys	Glu	Cys	Val	Ala	Pro	His	Met	Ala	Asp	Ile	Gly	Ser	Val	Thr
				165					170					175	
Ser	Ser	Ala	Ser	Gly	Pro	Ala	Gln	Ser	Val	Ala	Lys	Met	Phe	Arg	Ala
			180					185					190		
Val	Glu	Glu	Gly	Leu	Thr	Tyr	Lys	Phe	His	Ala	Ala	Trp	Ser	Ser	Val
		195					200					205			
Leu	Gln	Leu	Leu	Cys	Val	Phe	Phe	Glu	Ala	Cys	Gly	Arg	Gln	Ala	His
	210					215					220				
Pro	Val	Met	Arg	Lys	Cys	Leu	Gln	Ser	Leu	Cys	Asp	Leu	Arg	Leu	Ser
225					230					235					240
Pro	His	Phe	Pro	His	Thr	Ala	Ala	Leu	Asp	Gln	Ala	Val	Gly	Ala	Ala
				245					250					255	
Val	Thr	Ser	Met	Gly	Pro	Glu	Val	Val	l.eu	Gln	Ala	Val	Pro	Leu	Glu
			260					265					270		
He	Asp	Gly	Ser	Glu	Glu	Thr	Leu	Asp	Phe	Pro	Arg	Ser	Trp	Leu	Leu
		275					280					285			
Pro	Val	lle	Arg	Asp	His	Val	Gln	Glu	Thr	Arg	Leu	Gly	Phe	Phe	Thr
	290					295					300				
Thr	Tyr	Phe	Leu	Pro	Leu	Ala	Asn	Thr	Leu	Lys	Ser	Lys	Ala	Met	Asp
305					310					315					320

Leu	Ala	Gln	Ala	Gly	Ser	Thr	Val	Glu	Ser	Lys	Ile	Tyr	Asp	Thr	Leu
				325					330					335	
Gln	Trp	Gln	Met	Trp	Thr	Leu	Leu	Pro	Gly	Phe	Cys	Thr	Arg	Pro	Thr
			340					345					350		
Asp	Val	Ala	He	Ser	Phe	Lys	Gly	Leu	Ala	Arg	Thr	Leu	Gly	Met	Ala
		355					360					365			
Ile	Ser	Glu	Arg	Pro	Asp	Leu	Arg	Val	Thr	Val	Cys	Gln	Val	Leu	Arg
	370					375					380				
Thr	Leu	Ile	Thr	Lys	Gly	Cys	Gln	Ala	Glu	Ala	Asp	Arg	Ala	Glu	Val
385					390					395					400
Ser	Arg	Phe	Ala	Lys	Asn	Phe	Leu	Pro	He	Leu	Phe	Asn	Leu	Tyr	G1 y
				405					410					415	
Gln	Pro	Val	Ala	Ala	Gly	Asp	Thr	Pro	Ala	Pro	Arg	Arg	Ala	Val	Leu
			420					425					430		
Glu	Thr	lle	Arg	Thr	Tyr	Leu	Thr	He	Thr	Asp	Thr	Gln	Leu	Val	Asn
		435					440					445			
Ser	Leu	Leu	Glu	Lys	Ala	Ser	Glu	Lys	Val	Leu	Asp	Pro	Ala	Ser	Ser
	450					455					460				
Asp	Phe	Thr	Arg	Leu	Ser	Val	Leu	Asp	Leu	Val	Val	Ala	Leu	Ala	Pro
465					470					475					480
Cys	Ala	Asp	Glu	Ala	Ala	He	Ser	Lys	Leu	Tyr	Ser	Thr	lle	Arg	Pro
				485					490					495	
Tyr	Leu	Glu	Ser	Lys	Ala	His	Gly	Val	Gln	Lys	Lys	Ala	Tyr	Arg	Val
			500					505					510		
Leu	Glu	Glu	Val	Cys	Ala	Ser	Pro	G1n	Gly	Pro	Gly	Ala	Leu	Phe	Val
		515					520					525			
Gln	Ser	His	Leu	Glu	Asp	Leu	Lys	Lys	Thr	Leu	Leu	Asp	Ser	Leu	Arg
	530					535					540				
Ser	Thr	Ser	Ser	Pro	Ala	Lys	Arg	Pro	Arg	Leu	Lys	Cys	Leu	Leu	His
545					550					555					560
He	Val	Arg	Lys	Leu	Ser	Ala	Glu	His	Lys	Glu	Phe	He	Thr	Ala	Leu
				565					570					575	
lle	Pro	Glu		lle	Leu	Cys	Thr		Glu	Val	Ser	Val		Ala	Arg
			580					585					590		
Lys	Asn		Phe	Ala	Leu	Leu		Glu	Met	Gly	His			Leu	Arg
		595					600					605			

Phe	Gly	Ser	Asn	Gln	Glu	Glu	Ala	Leu	Gln	Cys	Tyr	Leu	Val	Leu	He
	610					615					620				
Tyr	Pro	Gly	Leu	Val	Gly	Ala	Val	Thr	Met	Val	Ser	Cys	Ser	lle	Leu
625					630					635					640
Ala	Leu	Thr	His	Leu	Leu	Phe	Glu	Phe	Lys	Gly	Leu	Met	G1 y	Thr	Ser
				645					650					655	
Thr	Val	Glu	Gln	Leu	Leu	Glu	Asn	Val	Cys	Leu	Leu	Leu	Ala	Ser	Arg
			660					665					670		
Thr	Arg	Asp	Val	Val	Lys	Ser	Ala	Leu	Gly	Phe	He	Lys	Val	Ala	Val
		675					680					685			
Thr	Val	Met	Asp	Val	Ala	His	Leu	Ala	Lys	His	Val	Gln	Leu	Val	Met
	690					695					700				
Glu	Ala	lle	Gly	Lys	Leu	Ser	Asp	Asp	Met	Arg	Arg	His	Phe	Arg	Met
705					710					715					720
Lys	Leu	Arg	Asn	Leu	Phe	Thr	Lys	Phe	He	Arg	Lys	Phe	G1 y	Phe	Glu
				725					730					735	
Leu	Val	Lys	Arg	Leu	Leu	Pro	Glu	Glu	Tyr	His	Arg	Va]	Leu	Val	Asn
			740					745					750		
Ile	Arg	Lys	Ala	Glu	Ala	Arg	Ala	Lys	Arg	His	Arg	Ala	Leu	Ser	Gln
		755					760					765			
Ala	Ala	Val	Glu	Pro	Ala										
	770					775					780				
Gln	Gly	Lys	Gly	Asp	Ser	lle	Glu	Glu	lle	Leu	Ala	Asp	Ser	Glu	Asp
785					790					795					800
Glu	Glu	Asp	Asn	Glu	Glu	Glu	Glu	Arg	Ser	Arg	Gly	Lys	Glu	Gln	Arg
				805					810					815	
Lys	Leu	Ala	Arg	Gln	Arg	Ser	Arg	Ala	Trp	Leu	Lys	Glu	Gly	Gly	Gly
			820					825					830		
Asp	Glu	Pro	Leu	Asn	Phe	Leu	Asp	Pro	Lys	Va]	Ala	Gln	Arg	Val	Leu
		835					840					845			
Ala	Thr	Gln	Pro	Gly	Pro	Gly	Arg	Gly	Arg	Lys	Lys	Asp	His	Gly	Phe
	850					855					860				
Lys	Val	Ser	Ala	Asp	Gly	Arg	Leu	lle	He	Arg	Glu	Glu	Ala	Asp	Gly
865					870					875					880
Asn	Lys	Met	Glu	G1u	Glu	Glu	Gly	Ala	Lys	Gly	Glu	Asp	Glu	Glu	Met
				885					890					895	

Ala Asp Pro Met Glu Asp Val Ile Ile Arg Asn Lys Lys His Gln Lys 900 905 Leu Lys His Gln Lys Glu Ala Glu Glu Glu Glu Leu Glu 11e Pro Pro 915 920 925 Gln Tyr Gln Ala Gly Gly Ser Gly Ile His Arg Pro Val Ala Lys Lys 935 Ala Met Pro Gly Ala Glu Tyr Lys Ala Lys Lys Ala Lys Gly Asp Val 950 955 Lys Lys Gly Arg Pro Asp Pro Tyr Ala Tyr Ile Pro Leu Asn Arg 970 965 Ser Lys Leu Asn Arg Arg Lys Lys Met Lys Leu Gln Gly Gln Phe Lys 985 980 Gly Leu Val Lys Ala Ala Gln Arg Gly Ser Gln Val Gly His Lys Asn 995 1000 1005 Arg Arg Lys Asp Arg Arg Pro 1010 1015

<210> 4793

<211> 445

<212> PRT

<213> Homo sapiens

<400> 4793

 Met
 Lys
 Asn
 Pro
 Glu
 Ala
 Gln
 Asp
 Val
 Ser
 Val
 Ser
 Gln
 Gly
 Phe

 1
 5
 5
 10
 10
 10
 15
 15
 15

 Arg
 Met
 Leu
 Phe
 Tyr
 Thr
 Met
 Lys
 Pro
 Ser
 Glu
 Thr
 Ser
 Phe
 Gln
 Thr
 Thr
 Asp
 Tyr
 Val
 Lys
 Lys
 Ala
 Thr
 Pro
 Phe
 Ph

85

90

Tyr	Ile	Tyr	Пe	Trp	Glu	Asn	Tyr	Arg	Leu	Phe	Asn	Leu	Pro	Trp	Asp
			100					105					110		
Ser	Pro	Trp	Thr	Trp	Tyr	Ser	Ala	Phe	Leu	Gly	Val	Asp	Phe	Gly	Tyr
		115					120					125			
Tyr	Trp	Phe	His	Arg	Met	Ala	His	Glu	Val	Asn	He	Met	Trp	Ala	Gly
	130					135					140				
His	Gln	Thr	His	His	Ser	Ser	Glu	Asp	Tyr	Asn	Leu	Ser	Thr	Ala	Leu
145					150					155					160
Arg	Gln	Ser	Val	Leu	Gln	He	Tyr	Thr	Ser	Trp	Ile	Phe	Tyr	Ser	Pro
				165					170					175	
Leu	Ala	Leu	Phe	Ile	Pro	Pro	Ser	Val	Tyr	Ala	Val	His	Leu	Gln	Phe
			180					185					190		
Asn	Leu		Tyr	Gln	Phe	Trp		His	Thr	Glu	Val		Asn	Asn	Leu
		195					200					205			
Gly		Leu	Glu	Leu	lle		Asn	Thr	Pro	Ser		His	Arg	Val	His
	210				_	215					220				
	Gly	Arg	Asn	Arg		Cys	He	Asp	Lys		Tyr	Ala	Gly	Val	
225		_		_	230					235					240
He	He	Trp	Asp		He	Phe	G1 y	Thr		Glu	Ala	Glu	Asn		Lys
		æ	61	245	m.		n	T 1	250	m)	TNI.	0.1	D	255	
Val	Val	lyr		Leu	lhr	H1S	Pro		Asn	Ihr	Phe	Glu	Pro	He	Lys
17. 3	6.1	DI	260		,	101	C	265	T	TC1	TI	DI	270 T	4.1	arı.
Val	GIn		HIS	HIS	Leu	Phe		11e	Trp	Ihr	Ihr		Trp	Ala	Inr
D	C1	275	DI	Λ	1	DL.	280	V = 1	11.	DI	1	285	D	C1	т
Pro		rne	rne	ASN	Lys		ser	vaj	116	rne		GIŸ	Pro	GIY	ırp
C1	290 Pro	C1	lvo	Dno	Ana	295	C1.,	Lou	Com	C1.,	300	110	Dno	C1	Vo.1
305	110	Gry	Lys	110	310	Leu	Gly	rea	261	315	Olu	116	Pro	Glu	320
	C1v	Lve	Glu	Val		Pho	Sor	Sor	Sor		Sor	Gln.	Leu	Lou	
1111	Uly	Lys	010	325	110	1116	261	261	330	261	361	OIII	Leu	335	1. y 3
ماا	Tyr	Thr	Val		Gln	Phe	Ala	Lau		Leu	Ala	Phe	Tyr		Glu
110		1111	340	101	0111	THE	MIG	345	MC C	Leu	Mid	THE	350	010	Olu
Thr	Phe	Ala		Thr	Ala	Ala	Leu		Gln	Val	Thr	leu	Leu	Leu	Aro
	,	355	чор	* 113		,,,,,,,	360	001	0.11			365	.500	200	*** 8
Val	Cvs		Πe	He	Leu	Thr		Thr	Ser	Πe	Glv		Leu	Len	Asn
	370					375					380				

<210> 4794

<211> 735

<212> PRT

130

145

<213> Homo sapiens

<400> 4794 Met Thr Cys Leu Ile Cys Ile Ala Ser Val Lys Arg Asn Gln Ala Val 10 Trp Ser Cys Ser Gly Cys Phe Cys Ile Phe His Met Pro Cys Ile Gln 25 20 30 Lys Trp Ala Lys Asp Ser Gln Phe Leu Val Ser Ser Val Thr Asp Asp 40 Asp Phe Gly Lys Lys Asp Cys Pro Trp Pro Cys Pro Lys Cys Arg Phe 50 55 60 Glu Tyr Lys Arg Ser Glu Thr Pro Ser Arg Tyr Tyr Cys Tyr Cys Gly 70 75 Lys Val Glu Asp Pro Pro Leu Asp Pro Trp Leu Val Pro His Ser Cys 90 Gly Gln Val Cys Glu Arg Glu Phe Lys Pro Pro Cys Gly His Lys Cys 100 105 110 Leu Leu Cys His Pro Gly Pro Cys Pro Pro Cys Pro Lys Met Val 120

Thr Thr Cys Tyr Cys Lys Lys Ala Lys Pro Ile Pro Arg Arg Cys

Ser Ala Lys Glu Trp Ser Cys Gln Leu Pro Cys Gly Gln Lys Leu Leu

155

160

135

Cys	Gly	Gln	llis	Lys	Cys	Glu	Asn	Pro	Cys	His	Ala	Gly	Ser	Cys	Gln
				165					170					175	
Pro	Cys	Pro	Arg	Val	Ser	Arg	Gln	Lys	Cys	Val	Cys	Gly	Lys	Lys	Val
			180					185					190		
Ala	Glu	Arg	Ser	Cys	Ala	Ser	Pro	Leu	Trp	His	Cys	Asp	Gln	Val	Cys
		195		,			200					205			
Gly	Lys	Thr	Leu	Pro	Cys	Gly	Asn	His	Thr	Cys	Glu	Gln	Val	Cys	His
	210					215					220				
Val	Gly	Ala	Cys	Gly	Glu	Cys	Pro	Arg	Ser	Gly	Lys	Arg	Phe	Cys	Pro
225					230					235					240
Cys	Gln	Lys	Ser	Lys	Phe	Ser	Leu	Pro	Cys	Thr	Glu	Asp	Val	Pro	Thr
				245					250					255	
Cys	Gly	Asp	Ser	Cys	Asp	Lys	Val	Leu	Glu	Cys	Gly	lle	His	Arg	Cys
			260					265					270		
Ser	Gln	Arg	Cys	His	Arg	Gly	Pro	Cys	Glu	Thr	Cys	Arg	Gln	Glu	Val
		275					280					285			
Glu	Lys	His	Cys	Arg	Cys	Gly	Lys	His	Thr	Lys	Arg	Met	Pro	Cys	His
	290					295					300				
Lys	Pro	Tyr	Leu	Cys	Glu	Thr	Lys	Cys	Val	Lys	Met	Arg	Asp	Cys	Gln
305					310					315					320
Lys	His	Gln	Cys	Arg	Arg	Lys	Cys	Cys	Pro	Gly	Asn	Cys	Pro	Pro	Cys
				325					330					335	
Asp	Gln	Asn	Cys	Gly	Arg	Thr	Leu	Gly	Cys	Arg	Asn	His	Lys	Cys	Pro
			340					345					350		
Ser	Val	Cys	His	Arg	Gly	Ser	Cys	Tyr	Pro	Cys	Pro	Glu	Thr	Val	Asp
		355					360					365			
Val	Lys	Cys	Asn	Cys	Gly	Asn	Thr	Lys	Val	Thr	Val	Pro	Cys	Gly	Arg
	370					375					380				
Glu	Arg	Thr	Thr	Arg	Pro	Pro	Lys	Cys	Lys	Glu	Gln	Cys	Ser	Arg	Pro
385					390					395					400
Pro	Thr	Cys	His	His	Thr	Ser	Gln	Glu	Lys	His	Arg	Cys	His	Phe	G1 y
				405					410					415	
Ser	Cys	Pro	Pro	Cys	His	GIn	Pro	Cys	Gln	Lys	Val	Leu	Glu	Lys	Cys
			420					425					430		
Gly	His	Leu	Cys	Pro	Ala	Pro	Cys	His	Asp	Gln	Ala	Leu	He	Lys	Gln
		435					440					445			

Thr	Gly	Arg	His	Gln	Pro	Thr	Gly	Pro	Trp	Glu	Gln	Pro	Ser	Glu	Pro
	450					455					460				
Ala	Phe	lle	G1n	Thr	Ala	Leu	Pro	Cys	Pro	Pro	Cys	Gln	Val	Pro	lle
465					470					475					480
Pro	Met	Glu	Cys	Leu	Gly	Lys	His	Glu	Val	Ser	Pro	Leu	Pro	Cys	His
				485					490					495	
Ala	Val	Gly	Pro	Tyr	Ser	Cys	Lys	Arg	Val	Cys	Gly	Arg	lle	Leu	Asp
			500					505					510		
Cys	Gln	Asn	,His	Thr	Cys	Met	Lys	Glu	Cys	His	Lys	Val	Thr	Lys	Thr
		515					520					525			
Asp	Gly	Cys	Thr	Gly	Lys	Asn	Lys	Ala	Gly	Pro	Glu	Cys	Leu	His	Cys
	530					535					540				
Glu	Glu	G1 y	Cys	Ser	Lys	Ser	Arg	Pro	Leu	Gly	Cys	Leu	His	Pro	Cys
545					550					555					560
Ile	Leu	Arg	Cys	His	Pro	Gly	Glu	Cys	Pro	Pro	Cys	Val	Gln	Met	Leu
				565					570					575	
Arg	lle	Lys	Cys	llis	Cys	Lys	lle	Thr	Ser	Leu	Tyr	Val	Glu	Cys	Arg
			580					585					590		
Lys	lle	Thr	Thr	Ala	Asp	Val	Asn	Glu	Lys	Asn	Leu	Leu	Ser	Cys	Cys
		595					600					605			
Lys	Asn	Gln	Cys	Pro	Lys	Glu	Leu	Pro	Cys	Gly	His	Arg	Cys	Lys	Glu
	610					615					620				
Met	Cys	His	Pro	Gly	Glu	Cys	Pro	Phe	Asn	Cys	Asn	Gln	Lys	Val	Lys
625					630					635					640
Leu	Arg	Cys	Pro	Cys	Lys	Arg	He	Lys	Lys	Glu	Leu	Gln	Cys	Asn	Lys
				645					650					655	
Val	Arg	Glu	Asn	Gln	Val	Ser	İle	Glu	Cys	Asp	Thr	Thr	Cys	Lys	Glu
			660					665					670		
Met	Lys	Arg	Lys	Ala	Ser	Glu	lle	Lys	Glu	Ala	Glu	Ala	Lys	Ala	Ala
		675					680					685			
Leu	Glu	Glu	Glu	Lys	Arg	Arg	Gln	Gln	Ala	Glu	Leu	Glu	Ala	Phe	Glu
	690					695					700				
Asn	Arg	Leu	Lys	Gly	Arg	Arg	Lys	Lys	Asn	Arg	Lys	Arg	Asp	Glu	Val
705					710					715					720
Ala	Val	Glu	Leu	Ser	Leu	Trp	Gln	Lys	His	Lys	Tyr	Tyr	Leu	lle	
				725					730					735	

```
<211> 578
<212> PRT
<213> Homo sapiens
<400> 4795
Met Pro Gly Pro His Leu Ser Leu Ser Ala Ile Ser Met Ser Leu Ser
                  5
                                     10
                                                          15
Leu His Leu Phe Leu Leu Pro Leu Ser Leu Tyr Phe Phe Ser Ser Leu
                                 25
Leu His Phe Gln Ser Leu Leu Leu Cys Leu Leu Val Ser Arg Phe Cys
         35
                             40
                                                  45
Val Ser lle Ser Leu Thr Cys lle Ser Met Ser Ser Phe Val Leu Phe
                         55
Ser Met Ser Leu Gly Arg His His Leu Leu Pro Thr Lys Ser Pro Tyr
                     70
                                         75
Pro His Ala Tyr Thr Ser Leu Leu Ser Leu Pro Gln Phe Asn Pro Glu
                 85
                                     90
                                                          95
Leu Val Leu Val Ser Ala Gly Phe Asp Ala Ala Arg Gly Asp Pro Leu
            100
                                105
Gly Gly Cys Gln Val Ser Pro Glu Gly Tyr Ala His Leu Thr His Leu
        115
                            120
                                                 125
Leu Met Gly Leu Ala Ser Gly Arg lle lle Leu Ile Leu Glu Gly Gly
                        135
Tyr Asn Leu Thr Ser lle Ser Glu Ser Met Ala Ala Cys Thr Arg Ser
                    150
                                        155
Leu Leu Gly Asp Pro Pro Pro Leu Leu Thr Leu Pro Arg Pro Pro Leu
                                     170
                165
                                                         175
Ser Gly Ala Leu Ala Ser lle Thr Glu Thr lle Gln Val His Arg Arg
                                185
Tyr Trp Arg Ser Leu Arg Val Met Lys Val Glu Asp Arg Glu Gly Pro
        195
                            200
                                                 205
Ser Ser Ser Lys Leu Val Thr Lys Lys Ala Pro Gln Pro Ala Lys Pro
    210
                        215
                                             220
```

<210> 4795

Arg	Leu	Ala	Glu	Arg	Met	Thr	Thr	Arg	Glu	Lys	Lys	Val	Leu	Glu	Ala
225					230					235					240
G1y	Met	Gly	Lys	Val	Thr	Ser	Ala	Ser	Phe	Gly	Glu	Glu	Ser	Thr	Pro
				245					250					255	
Gly	Gln	Thr	Asn	Ser	Glu	Thr	Ala	Val	Val	Ala	Leu	Thr	Gln	Asp	Gln
			260					265					270		
Pro	Ser	Glu	Ala	Ala	Thr	G1 y	Gly	Ala	Thr	Leu	Ala	Gln	Thr	He	Ser
		275					280					285			
Glu	Ala	Ala	Ile	Gly	Gly	Ala	Met	Leu	Gly	Gln	Thr	Thr	Ser	Glu	Glu
	290					295					300				
Ala	Val	Gly	Gly	Ala	Thr	Pro	Asp	Gln	Thr	Thr	Ser	Glu	Glu	Thr	Val
305					310					315					320
Gly	Gly	Ala	He	Leu	Asp	Gln	Thr	Thr	Ser	Glu	Asp	Ala	Val	Gly	Gly
				325					330					335	
Ala	Thr	Leu	Gly	Gln	Thr	Thr	Ser	Glu	Glu	Ala	Val	Gly	Gly	Ala	Thr
			340					345					350		
Leu	Ala	Gln	Thr	Thr	Ser	Glu	Ala	Ala	Met	Glu	Gly	Ala	Thr	Leu	Asp
		355					360					365			
Gln	Thr	Thr	Ser	Glu	Glu	Ala	Pro	Gly	Gly	Thr	Glu	Leu	Ile	Gln	Thr
	370					375					380				
Pro	Leu	Ala	Ser	Ser	Thr	Asp	His	Gln	Thr	Pro	Pro	Thr	Ser	Pro	Val
385					390					395					400
Gln	Gly	Thr	Thr	Pro	Gln	He	Ser	Pro	Ser	Thr	Leu	11e	Gly	Ser	Leu
				405					410					415	
Arg	Thr	Leu	Glu	Leu	Gly	Ser	Glu	Ser	Gln	Gly	Ala	Ser	Glu	Ser	Gln
			420					425					430		
Ala	Pro	Gly	Glu	Glu	Asn	Leu	Leu	Gly	Glu	Ala	Ala	Gly	Gly	Gln	Asp
		435					440					445			
Met	Ala	Asp	Ser	Met	Leu	Met	Gln	Gly	Ser	Arg	Gly	Leu	Thr	Asp	Gln
	450					455					460				
Ala	lle	Phe	Tyr	Ala	Val	Thr	Pro	Leu	Pro	Trp	Cys	Pro	His	Leu	Val
465					470					475					480
Ala	Val	Cys	Pro	He	Pro	Ala	Ala	Gly	Leu	Asp	Val	Thr	Gln	Pro	Cys
				485					490					495	
Gly	Asp	Cys	Gly	Thr	He	G]n	Glu	Asn	Trp	Val	Cys	Leu	Ser	Cys	Tyr
			500					505					510		

Gln Val Tyr Cys Gly Arg Tyr Ile Asn Gly His Met Leu Gln His His
515 520 525

Gly Asn Ser Gly His Pro Leu Val Leu Ser Tyr Ile Asp Leu Ser Ala

530 535 540

Trp Cys Tyr Tyr Cys Gln Ala Tyr Val His His Gln Ala Leu Leu Asp 545 550 555 560

Val Lys Asn 11e Ala His Gln Asn Lys Phe Gly Glu Asp Met Pro His 565 570 575

Pro His

<210> 4796

<211> 108

<212> PRT

<213> Homo sapiens

<400> 4796

Met Gly Thr Asp Glu Asp Cys Tyr Gln Glu Arg Gly Asp Trp Asp Gln

1 5 10 15

Glu Ser Gly Phe Gly Arg Glu Glu Pro Arg Leu Leu Phe Trp Met Cys
20 25 30

Gln Asp Arg Ser Ala Phe Glu Lys Ser Glu Leu Trp Tyr Gln Ile Ala 35 40 45

Ser Trp Jle Cys Glu Thr Gly Val Trp Gly Lys Glu Arg Pro Trp Leu 50 55 60

Lys Asn Val Ser Phe Gly Asn Val Asn lle Lys Pro Gln Asp Trp Val 65 70 75 80

Thr Trp Pro Ser Arg Gly Val Gly Thr Glu Lys Arg Pro Trp Pro Glu

85 90 95

Pro Trp Gly His Leu His Leu Gly Glu Lys Gly Met

<211> 142 <212> PRT <213> Homo sapiens <400> 4797 Met Ala Pro Val Lys IIe Ser His Val Val Ser Phe Ser Ser Gln Asp 10 Pro Lys Tyr Pro Val Glu Asn Leu Leu Asn Pro Asp Ser Pro Arg Arg 20 25 30 Pro Trp Leu Gly Cys Pro Gln Asp Lys Ser Gly Gln Leu Lys Val Glu 35 40 Leu Gln Leu Glu Arg Ala Val Pro Thr Gly Tyr Ile Asp Val Gly Asn 50 55 60 Cys Gly Cys Ala Phe Leu Gln Ile Asp Val Gly His Ser Ser Trp Pro 65 70 75 Leu Asp Arg Pro Phe Ile Thr Leu Leu Pro Ala Thr Thr Leu Met Ser 90 Leu Thr Asp Ser Lys Gln Gly Lys Asn Arg Ser Gly Val Arg Met Phe 100 105 110 Lys Asp Gly Lys Glu Gly Lys Ser Arg Lys Asp Gly Gly Leu Tyr 115 120 Glu Lys Gln Arg Cys Ser Thr Lys Glu Asp Cys Glu Cys Tyr 130 135 140 <210> 4798 <211> 153 <212> PRT <213> Homo sapiens <400> 4798 Met Tyr Glu lle Tyr Val Glu Thr Cys Gly Gln Asn Thr Glu Asn Gln Val Asn Pro Ala Thr Phe Gly Lys Leu Val Arg Leu Val Phe Pro Asp 20 25 30

Leu Gly Thr Arg Trp Leu Gly Thr Arg Gly Ser Ala Arg Tyr His Tyr

40 Asp Gly 11e Cys 11e Lys Lys Ser Ser Phe Phe Tyr Ala Gln Tyr Cys 55 Cys Leu lle Gly Glu Lys Arg Tyr His Ser Gly Asp Ala lle Ala Phe 75 70 Glu Lys Ser Thr Asn Tyr Asn Ser Ile Ile Gln Gln Glu Ala Thr Cys 90 85 Glu Asp His Ser Pro Met Lys Thr Asp Pro Val Gly Ser Pro Leu Ser 100 105 110 Glu Phe Arg Arg Cys Pro Phe Leu Glu Gln Glu Leu Ala Lys Lys Tyr 115 120 125 Ser Cys Asn Met Met Ala Phe Leu Ala Asp Glu Tyr Cys Asn Tyr Cys 135 140 Arg Asp lle Leu Arg Asn Val Arg Asn 145 150

<210> 4799

<211> 787

<212> PRT

<213> Homo sapiens

<400> 4799

Met Glu Gly Ala Gly Glu Asn Ala Pro Glu Ser Ser Ser Ser Ala Pro

1 5 10 15

Gly Ser Glu Glu Ser Ala Arg Asp Pro Gln Val Pro Pro Pro Glu Glu 20 25 30

Glu Ser Gly Asp Cys Ala Arg Ser Leu Glu Ala Val Pro Lys Lys Leu 35 40. 45

Cys Gly Tyr Leu Ser Lys Phe Gly Gly Lys Gly Pro 11e Arg Gly Trp
50 55 60

Lys Ser Arg Trp Phe Phe Tyr Asp Glu Arg Lys Cys Gln Leu Tyr Tyr
65 70 75 80

Ser Arg Thr Ala Gln Asp Ala Asn Pro Leu Asp Ser Val Asp Leu Ser 85 90 95

Ser Ala Val Phe Asp Cys Lys Ala Asp Ala Glu Glu Gly Ile Phe Glu

			100					105					110		
He	Lys	Thr	Pro	Ser	Arg	Val	He	Thr	Leu	Lys	Ala	Ala	Thr	Lys	Gln
		115					120					125			
Ala	Met	Leu	Tyr	Trp	Leu	Gln	Gln	Leu	Gln	Met	Lys	Arg	Trp	Glu	Phe
	130					135					140				
His	Asn	Ser	Pro	Pro	Ala	Pro	Pro	Ala	Thr	Pro	Asp	Ala	Ala	Leu	Ala
145					150					155					160
Gly	Asn	Gly	Pro	Val	Leu	His	Leu	Glu	Leu	Gly	Gln	Glu	Glu	Ala	Glu
				165					170					175	
Leu	Glu	Glu	Phe	Leu	Cys	Pro	Val	Lys	Thr	Pro	Pro	G1y	Leu	Val	G1 y
			180					185					190		
Val	Ala	Ala	Ala	Leu	Gln	Pro	Phe	Pro	Ala	Leu	Gln	Asn	11e	Ser	Leu
		195					200					205			
Lys	His	Leu	Gly	Thr	Glu	He	Glń	Asn	Thr	Met	His	Asn	Пе	Arg	Gly
	210					215					220				
Asn	Lys	Gln	Ala	Gln	Gly	Thr	Gly	His	Glu	Pro	Pro	Gl y	Glu	Asp	Ser
225					230					235					240
Pro	Gln	Ser	Gly	Glu	Pro	Gln	Arg	Glu	Glu	Gln	Pro	Leu	Ala	Ser	Asp
				245					250					255	
Ala	Ser	Thr	Pro	Gly	Arg	Glu	Pro	Glu	Asp	Ser	Pro	Lys	Pro	Ala	Pro
			260					265					270		
Lys	Pro	Ser	Leu	Thr	lle	Ser	Phe	Ala	Gln	Lys	Ala	Lys	Arg	Gln	Asn
		275					280					285			
Asn	Thr	Phe	Pro	Phe	Phe	Ser	Glu	G1 y	He	Thr	Arg	Asn	Arg	Thr	Ala
	290					295					300				
Gln	Glu	Lys	Val	Ala	Ala	Leu	Glu	Gln	Gln	Val	Leu	Met	Leu	Thr	Lys
305					310					315					320
Glu	Leu	Lys	Ser		Lys	Glu	Leu	Val	Lys	He	Leu	His	Lys		Leu
				325					330					335	
G1u	Ala	Ala		Gln	Glu	Lys	Arg		Ser	Ser	Ala	Tyr		Ala	Ala
			340					345					350		
Ala	Glu		Lys	Asp	Arg	Leu		Leu	Val	Arg	His		Val	Arg	Gln
		355					360					365			
He		Glu	Leu	Gly	Arg	Arg	Val	Glu	Ala	Leu		Gln	Glu	Arg	Glu
_	370					375					380		۵.	٥.	
Ser	Leu	Ala	His	Thr	Ala	Ser	Leu	Arg	Glu	Gln	GIn	Val	GIn	Glu	Leu

;	385					390					395					400
(Gln	G1n	His	Val	Gln	Leu	Leu	Met	Asp	Lys	Asn	His	Ala	Lys	Gln	Gln
					405					410					415	
١	Val	He	Cys	Lys	Leu	Ser	Glu	Lys	Val	Thr	Gln	Asp	Phe	Thr	His	Pro
				420					425					430		
l	Pro	Asp	Gln	Ser	Pro	Leu	Arg	Pro	Asp	Ala	Ala	Asn	Arg	Asp	Phe	Leu
			435					440					445			
	Ser	G1n	Gln	Gly	Lys	Ile	Glu	His	Leu	Lys	Asp	Asp	Met	Glu	Ala	Tyr
		450					455					460				
1	Arg	Thr	Gln	Asn	Cys	Phe	Leu	Asn	Ser	Glu	Ile	His	G1n	Val	Thr	Lys
,	465					470					475					480
	He	Trp	Arg	Lys	Val	Ala	Glu	Lys	Glu	Lys	Ala	Leu	Leu	Thr	Lys	Cys
					485					490					495	
ı	Ala	Tyr	Leu	Gln	Ala	Arg	Asn	Cys	Gln	Val	G1u	Ser	Lys	Tyr	Leu	Ala
				500					505					510		
(Gly	Leu	Arg	Arg	Leu	Gln	Glu	Ala	Leu	Gly	Asp	Glu	Ala	Ser	Glu	Cys
			515					520					525			
,	Ser	Glu	Leu	Leu	Arg	Gln	Leu	Val	Gln	Glu	Ala	Leu	Gln	Trp	Glu	Ala
		530					535					540				
(Gly	Glu	Ala	Ser	Ser	Asp	Ser	He	Glu	Leu	Ser	Pro	He	Ser	Lys	Tyr
į	545					550					555					560
ı	Asp	Glu	Tyr	Gly	Phe	Leu	Thr	Val	Pro	Asp	Tyr	Glu	Val	Glu	Asp	Leu
					565					570					575	
	Lys	Leu	Leu	Ala	Lys	lle	Gln	Ala	Leu	Glu	Ser	Arg	Ser	His	His	Leu
				580					585					590		•
]	Leu	Gly	Leu	Glu	Ala	Val	Asp	Arg	Pro	Leu	Arg	Glu	Arg	Trp	Val	Ala
			595					600					605			
]	Leu	Gly	Asp	Leu	Val	Pro	Ser	Ala	Glu	Leu	Lys	Gln	Leu	Leu	Arg	Ala
		610					615					620				
(Gly	Val	Pro	Arg	Glu	His	Arg	Pro	Arg	Val	Trp	Arg	Trp	Leu	Va]	His
(625					630					635					640
	Leu	Arg	Val	Gln	His	Leu	His	Thr	Pro	Gly	Cys	Tyr	Gln	G] u	Leu	Leu
					645					650					655	
;	Ser	Arg	Gly	Gln	Ala	Arg	Glu	His	Pro	Ala	Ala	Arg	Gln	11e	Glu	Leu
				660					665					670		
,	Asp	Leu	Asn	Arg	Thr	Phe	Pro	Asn	Asn	Lys	His	Phe	Thr	Cys	Gly	Ser

		675					680					685			
Trp	Ser	Ser	Leu	Arg	Gln	Ser	Thr	Trp	Gly	Gly	Gly	His	Pro	Gly	Ala
	690					695					700				
Glu	Leu	Cys	Pro	Arg	Ala	Val	Pro	Ala	Arg	Thr	Arg	Trp	Arg	Gly	Lys
705					710					715					720
Pro	Asp	Leu	Ala	Thr	Ser	Pro	Pro	His	Ser	Leu	Pro	His	Pro	Trp	Leu
				725					730					735	
Ala	Asp	Pro	Leu	Glu	Val	Arg	His	Gly	Pro	Val	Ala	Gln	Pro	Trp	Val
			740					745					750		
Ser	His	His	His	Val	Thr	Leu	Asp	Met	Ser	Leu	Pro	Leu	Ser	Gly	Pro
		755					760					765			
Gln	Phe	Pro	His	Trp	Asp	Ile	Val	Cys	Cys	Lys	Ala	Ile	Gly	Trp	Ala
	770					775					780				
Thr	Ser	Ser													
785															
<210)> 48	300													
<21	1> 22	21													
<212	2> PI	RT													
<213	3> Ho	omo :	sapi	ens											
)> 48														
Met	Ser	Gly	Glu	Ala	Thr	Val	Leu	Ala		His	Ala	Pro	Glu		Gln
1				5					10					15	
Glu	Gly	Leu	Leu	Val	Val	Lys	Val		Glu	Glu	Asn	Tyr		Leu	Asp
			20					25	_				30		
Gln	Asp		Gly	Leu	Gln	GJu		Pro	Trp	Ser	GIn		Val	Phe	Arg
		35			•••	~	40	•		2	are i	45			61
GIn		Phe	Arg	GIn	Phe		Tyr	Ser	Asp	Ser		Gly	Pro	Arg	GJU
	50					55			_		60			15	0.1
	Leu	Ser	Arg	Leu		Glu	Leu	Cys	Cys		Trp	Leu	Arg	Pro	
65		_	_	0.1	70					75			0.1	0.1	80
Val	His	Ser	Lys		GIn	He	Leu	Glu		Leu	Met	Leu	Glu		Phe
		**		85	0.1	0.1		0.1	90	an a	•		C.	95	
Leu	Ala	He	Leu	Pro	Glu	Glu	Leu	Gln	Ala	Trp	Leu	Arg	Glu	His	Arg

105 110 100 Pro Glu Asn Gly Glu Glu Ala Val Thr Met Leu Glu Glu Leu Glu Lys 120 125 Glu Leu Glu Glu Pro Arg Gln Gln Asp Thr Thr His Gly Gln Glu Met 135 140 Phe Trp Gln Glu Met Thr Ser Thr Gly Ala Leu Lys Ser Leu Ser Leu 155 150 Asn Ser Pro Val Gln Pro Leu Glu Asn Gln Cys Lys Thr Glu Thr Gln 170 175 165 Glu Ser Gln Ala Phe Gln Glu Arg Asp Gly Val Ser Leu Cys His Pro 180 185 Gly Trp Ser Ala Val Val Gln Pro Gln Leu Thr Ala Val Ala Leu Asn 200 Pro Trp Val Lys Val Ile Leu Leu Pro Gln Pro Pro Glu 210 215 220

<210> 4801

<211> 1043

<212> PRT

<213> Homo sapiens

<400> 4801

Met Ala Met Ala Leu Glu Leu Gln Ala Gln Ala Ser Pro Gln Pro Glu
1 5 10 15

Pro Glu Glu Leu Leu Ile Val Lys Leu Glu Glu Asp Ser Trp Gly Ser
20 25 30

Glu Ser Lys Leu Trp Glu Lys Asp Arg Gly Ser Val Ser Gly Pro Glu 35 40 45

Ala Ser Arg Gln Arg Phe Arg Gln Phe Gln Tyr Arg Asp Ala Ala Gly
50 55 60

Pro His Glu Ala Phe Ser Gln Leu Trp Ala Leu Cys Cys Arg Trp Leu
65 70 75 80

Arg Pro Glu lle Arg Leu Lys Glu Gln lle Leu Glu Leu Leu Val Leu 85 90 95

Glu Gln Phe Leu Thr lle Leu Pro Arg Glu Val Gln Thr Trp Val Gln

			100					105					110		
Ala	Arg	His	Pro	Glu	Ser	Gly	Glu	Glu	Ala	Val	Ala	Leu	Val	Glu	Asp
		115					120					125			
Trp	His	Arg	Glu	Thr	Arg	Thr	Ala	Gly	Gln	Ser	Gly	Leu	Glu	Leu	His
	130					135					140				
Thr	Glu	Glu	Thr	Arg	Pro	Leu	Lys	Thr	Gly	Glu	Glu	Ala	Gln	Ser	Phe
145					150					155					160
G1n	Leu	Gln	Pro	Val	Asp	Pro	Trp	Pro	Glu	Gly	Gln	Ser	Gln	Lys	Lys
				165					170					175	
Gly	Val	Lys	Asn	Thr	Cys	Pro	Asp	Leu	Pro	Asn	His	Leu	Asn	Ala	Glu
			180					185					190		
Val	Ala	Pro	Gln	Pro	Leu	Lys	Glu	Ser	Ala	Val	Leu	Thr	Pro	Arg	Val
		195					200					205			
Pro	Thr	Leu	Pro	Lys	Met	Gly	Ser	Val	Gly	Asp	Trp	Glu	Val	Thr	Ala
	210					215					220				
Glu	Ser	Gln	Glu	Ala	Leu	Gly	Pro	Gly	Lys	His	Ala	Glu	Lys	Glu	Leu
225					230					235					240
Cys	Lys	Asp	Pro	Pro	Gly	Asp	Asp	Cys	Gly	Asn	Ser	Va]	Cys	Leu	Gly
				245					250					255	
Val	Pro	Val	Ser	Lys	Pro	Ser	Asn	Thr	Ser	Glu	Lys	Glu	Gln	Gly	Pro
			260					265					270		
Glu	Phe	Trp	Gly	Leu	Ser	Leu	Ile	Asn	Ser	Gly	Lys	Arg	Ser	Thr	Ala
		275					280					285			
Asp		Ser	Leu	Asp	Asn	Glu	Pro	Ala	Gln	Ala	Leu	Thr	Trp	Arg	Asp
	290					295			_		300				
		Ala	Trp	Glu		Gln		Gln	Trp			Glu	Asp	Met	
305						0.1		0.1	61	315			5 1		320
Val	Ser	Gly	Val		Trp	Gly	Tyr	Glu		Thr	Lys	Thr	Phe		Ala
	,	0	61	325	р	101		<i>a</i> 1	330	,		Tr.	0	335	61
lle	Leu	Ser		Ser	Pro	Phe	Ser		Lys	Leu	Arg	Ihr		HIS	GIN
			340					345					350		
Asn	Arg	Gln	Val	Tyr	Arg	Ala	Ile	Ala	Glu	Gln	Leu	Arg	Ala	Arg	Gly
		355					360					365			
Phe	Leu	Arg	Thr	Leu	Glu	G]n	Cys	Arg	Tyr	Arg	Val	Lys	Asn	Leu	Leu
	370					375					380				

Arg	Asn	Tyr	Arg	Lys	Ala	Lys	Ser	Ser	His	Pro	Pro	Gly	Thr	Cys	Pro
385					390					395					400
Phe	Tyr	Glu	Glu	Leu	Glu	Ala	Leu	Val	Arg	Ala	Arg	Thr	Ala	lle	Arg
				405					410					415	
Ala	Thr	Asp	Gly	Pro	Gly	Glu	Ala	Val	Ala	Leu	Pro	Arg	Leu	Gly	Asp
			420					425					430		
Ser	Asp	Ala	Glu	Met	Asp	Glu	Gln	Glu	Glu	Gly	Gly	Trp	Asp	Pro	Glu
		435					440					445			
Glu	Met	Ala	Glu	Asp	Cys	Asn	Gly	Ala	Gly	Leu	Val	Asn	Val	Glu	Ser
	450					455					460				
Thr	Gln	Gly	Pro	Arg	He	Ala	Gly	Ala	Pro		Leu	Phe	Gln	Ser	
465					470					475					480
He	Ala	G]y	Val	His	Trp	Gly	Tyr	Glu		Thr	Lys	Ala	Phe		Ala
				485					490					495	
He	Leu	Ser	Glu	Ser	Pro	Phe	Ser		Lys	Leu	Arg	Thr		His	GIn
			500					505					510		0.1
Asn	Ser		Val	Tyr	Arg	Ala		Ala	Glu	Arg	Leu		Ala	Leu	Gly
		515	mı		0.1	0.1	520		æ		DI	525			
Phe		Arg	Thr	Leu	Glu		Cys	Arg	lyr	Arg		Lys	Asn	Leu	Leu
	530	т	Α.	1	41.	535	C	C	112 -	Dana	540	C1	Than	Cua	Dwo
	5er	ıyr	Arg	Lys		Lys	ser	ser	nis		PTO	Gry	1111	Cys	560
545	т	C1	C1	Lau	550	Com	Lau	Mot	Ana	555	Ana	410	A10	Vo.1	
rne	ryr	Gju	Glu	565	ASP	261	Leu	мес	570	MIA	Mg	MIA	nia	575	nı g
410	Mat	C1v	Thr		Ara	C1 o	Λla	Ala		Lou	Pro	Δra	Cvs		Gln
МІа	мес	U.I Y	580	vai	MIG	Olu	nia	585	Gry	Leu	110	MI g	590	Oly	OIII
Sor	Sor	Ala	Glu	Thr	Asn	Ala	Gln		Ala	Trn	Glv	Glu		Ala	Asn
001	001	595			пор	,,, d	600				01)	605			
Glu	Asn		Val	Lvs	Pro	Ser			Cvs	Pro	Lvs			Asp	Met
014	610	7110		13,10		615		200	0,0		620				
Glv		Glu	Met	Arg	His			Glu	Asp	Gln			Glu	Gln	Asp
625				J	630				•	635					640
	Phe	Glu	Gly	Leu			Λla	Leu	Ser	Lys	Cys	Pro	Thr	Glu	Ala
			·	645					650		-			655	
Val	Cys	Gln	Pro			Trp	Gly	Glu	Asp	Ser	Glu	Asn	Glu	Asn	Glu
	•		660			·	·	665					670		

Asp	Glu	Gly	Gln	Trp	Gly	Asn	Pro	Ser	Gln	Glu	Gln	Trp	Gln	Glu	Ser
		675					680					685			
Ser	Ser	Glu	Glu	Asp	Leu	Glu	Lys	Leu	lle	Asp	His	Gln	Gly	Leu	Tyr
	690					695					700				
Leu	Ala	Glu	Lys	Pro	Tyr	Lys	Cys	Asp	Thr	Cys	Met	Lys	Ser	Phe	Ser
705					710					715					720
Arg	Ser	Ser	His	Phe	lle	Ala	His	Gln	Arg	lle	His	Thr	Gly	Glu	Lys
				725					730					735	
Pro	Tyr	Lys	Arg	Leu	Glu	Cys	Gly	Lys	Asn	Phe	Ser	Asp	Arg	Ser	Asn
			740					745					750		
Leu	Asn	Thr	His	Gln	Arg	He	His	Thr	Gly	Glu	Lys	Pro	Tyr	Lys	Cys
		755					760					765			
Leu	Glu	Cys	Gly	Lys	Ser	Phe	Ser	Asp	His	Ser	Asn	Leu	lle	Thr	His
	770					775					780				
Gln	Arg	lle	His	Thr	Gly	Glu	Lys	Pro	Tyr	Lys	Cys	Gly	Glu	Cys	Trp
785					790					795					800
Lys	Ser	Phe	Asn	Gln	Ser	Ser	Asn	Leu	Leu	Lys	His	Gln	Arg	Ile	His
				805					810					815	
Leu	Gly	Gly	Asn	Pro	Asp	Gln	Cys	Ser	Glu	Pro	Gly	G1 y	Asn	Phe	Ala
			820					825					830		
Gln	Ser	Pro	Ser	Phe	Ser	Ala	His	Trp	Arg	Asn	Ser		Glu	Glu	Thr
		835					840					845			
Ala	Pro	Glu	Gln	Pro	Gln		He	Ser	Lys	Asp		Asn	Ser	Pro	Gly
	850					855					860				
Pro	His	Ser	Thr	Asn	Ser	Gly	Glu	Lys	Leu			Cys	Ser	Glu	
865					870					875					880
Gly	Arg	Ser	Phe		Lys	Ser	Ser	Ala			Ser	His	GIn		He
				885	_	_		_	890		_			895	D1
His	Thr	Gly		Lys	Pro	Tyr	GIu			GIu	Cys	Gly		Ser	Phe
			900	m.i				905			m		910	61	61
Ser	Lys			Thr	Leu	Ala			GIn	Arg	Thr			GIy	Glu
		915		0	1		920		,	•	151	925			C
Lys		Tyr	Lys	Cys	Val			Gly	Lys	Cys			61u	Arg	Ser
	930	77	T.		C1	935			Tì	C1	940		D-	т	1
		He	Ihr	H1S	Gln		val	H1S	Ihr			Lys	rro	ıyr	
945					950					955					960

Cys Leu Glu Cys Gly Lys Phe Phe Arg Asp Arg Ser Asn Leu Ile Thr 970 His Gln Arg lle His Thr Gly Glu Lys Pro Tyr Lys Cys Arg Glu Cys 980 985 990 Gly Lys Cys Phe Asn Gln Ser Ser Ser Leu lle Ile His Gln Arg Ile 1000 His Thr Gly Glu Lys Pro Tyr Lys Cys Thr Glu Cys Gly Lys Asp Phe 1015 1020 Asn Asn Ser Ser His Phe Ser Ala His Arg Arg Thr His Ala Gly Gly 1035 1040 1025 1030 Lys Ala Ser

<210> 4802

<211> 422

<212> PRT

<213> Homo sapiens

<400> 4802

Met Pro Leu Trp Glu Ala Pro Ser Asp His Pro Ala Asn Pro Pro Ala 10 Thr Leu Gln Gly His Thr Ser Leu Pro Gly Cys Gln Glu Glu Pro Ala 20 25 30 Arg Asp Pro Gln Ser Gly Leu Pro Gln Ile Thr Ser Glu Ser Ser Ser 40 Phe Ser Glu Gly Ser Leu Pro Ser Trp Ser Ser Gly Pro Ala Gly Ala 50 55 Lys Leu Asn Ala Scr His Glu Gly 11e Gly Scr Scr Scr Asp Gly Asn 65 Gly Asp Ser Lys Ala Ala Thr Glu Arg Val Val Ser Ala Met Asp Thr 85 90 Val Arg Arg Lys His Pro Glu IIe Thr Phe Tyr IIe Leu Val Lys Ala 110 100 105

lle Tyr Thr Leu Gly Tyr Ser Val Ser Leu Met Ser Leu Ala Thr Gly

		115					120					125			
Ser	Пe	lle	Leu	Cys	Leu	Phe	۸rg	Lys	Leu	His	Cys	Thr	Arg	Asn	Tyr
	130					135					140				
He	His	Leu	Asn	Leu	Phe	Leu	Ser	Phe	Пe	Leu	۸rg	Ala	Πe	Ser	Val
145					150					155					160
Leu	Val	Lys	Asp	Asp	Val	Leu	Tyr	Ser	Ser	Ser	Gly	Thr	Leu	His	Cys
				165					170					175	
Pro	Asp	Gln	Pro	Ser	Ser	Trp	Val	Gly	Cys	Lys	Leu	Ser	Leu	Val	Phe
			180					185					190		
Leu	G1n	Tyr	Cys	lle	Met	Ala	Asn	Phe	Phe	Trp	Leu	Leu	Val	Glu	Gly
		195					200					205			
Leu	Tyr	Leu	His	Thr	Leu	Leu	Val	Ala	Met	Leu	Pro	Pro	Arg	Arg	Cys
	210					215					220				
Phe	Leu	Ala	Tyr	Leu	Leu	He	Gly	Trp	Gly	Leu	Pro	Thr	Val	Cys	lle
225					230					235					240
G1 y	Ala	Trp	Thr	Ala	Ala	Arg	Leu	Tyr	Leu	Glu	Asp	Thr	Gly	Cys	Trp
				245					250					255	
Asp	Thr	Asn	Asp	His	Ser	Val	Pro		Trp	Val	Ile	Arg		Pro	He
			260					265					270		
Leu	lle		lle	He	Val	Asn		Val	Leu	Phe	lle		He	He	Arg
		275					280					285	-1		
He		Leu	Gln	Lys	Leu		Ser	Pro	Asp	Val		Gly	Asn	Asp	GIn
	290					295			m)		300			13	,
	GIn	Tyr	Lys	Arg	Leu	Ala	Lys	Ser	lhr		Leu	Leu	116	Pro	
305	C1	V 1	11:	т	310	W - 1	DL -	۸1.	V - I	315	Dece	11.	Can	11.	320
Phe	GIY	vai	HIS	-	Met	vai	Pne	АТа	330		Pro	116	ser	335	
Con	Luc	Tue	Cln	325		Dho	C1.	Lou			C1v	Sor	Pho		
sei	Lys	1 y 1	340	116	Leu	THE	Olu	345	Cys	Leu	Oly	361	350	0111	Oly
Lou	Val	Val		Val	Leu	Tyr	Cve		Lou	Asn	Sor	Glu		Gln	Cvs
Leu	101	355	MIG	1 (1)	Leu	1 y 1	360	1 110	LCG	11311	561	365		,	O y G
Glu	leu		Arσ	Lve	Trp	Arø		Arø	Cvs	Pro	Thr		Ser	Ala	Ser
Old	370	223	g	10,0	ıιρ	375		8	0,0		380		~03		
Arø		Tvr	Arø	Val	Cys		Ser	Ser	Phe	Ser		Asn	Glv	Ser	Glu
385			6		390					395			3		400
	Ala	Len	Gln	Phe	His	Arσ	G1 v	Ser	Ara			Ser	Phe	Leu	

405 410 415

Thr Glu Thr Ser Val 11e 420

<210> 4803

<211> 310

<212> PRT

<213> Homo sapiens

<400> 4803

Met Pro Ser Ser Lys Pro Gly Ser Pro Ala Pro Gln Asp Pro Ser Pro 1 5 10 15

Ser Cys Arg Thr Cys Ser Asn Gln Val Gln Pro Gly His Leu Leu Leu 20 25 30

Pro Leu Thr Lys Ala Ser Leu Thr Gly Pro Ser Pro Pro Pro Ala Lys
35 40 45

Leu Gly Arg Ala Gln Gly Ala Pro Ser Pro Lys Gly Pro Leu Glu Ile 50 55 60

Gly Trp Ser Pro Asp Ala Gln Pro Leu Arg Leu Gly Glu Thr Phe Leu
65 70 75 80

Pro Lys Gly Val Ala Gln Leu Ser Arg Ala Ser Ser Ser Gly His Lys
85 90 95

Val Gly Phe Leu Gln Pro Trp Pro Pro His Pro Leu Leu Pro Gln Glu 100 105 110

Arg Val Pro Arg His Gln Arg Gln Cys Cys Pro Arg Ala Gly Pro Ala 115 120 125

Phe Gly Cys Leu Arg Gly Arg Pro Gly Gly Trp Pro Ala Thr Gly Ser 130 135 140

Gly Gly Arg Gln Gln Gly Arg Lys Met Gly His Pro Pro Val Ser Pro 145 150 155 160

Ser Ala Pro Ala Pro Ala Gly Thr Thr Ala lle Pro Gly Leu lle Pro 165 170 175

Asp Leu Val Ala Gly Thr Pro Trp Pro Arg Trp Ala Leu Ile Ala Gly 180 185 190

Ala Leu Ala Ala Gly Val Leu Leu Val Ser Cys Leu Leu Cys Ala Ala

		195					200					205			
Cys	Cys	Cys	Cys	Arg	Arg	His	Arg	Lys	Lys	Pro	Arg	Asp	Lys	Glu	Ser
	210					215					220				
Val	Gly	Leu	Gly	Ser	Ala	Arg	Gly	Thr	Thr	Thr	Thr	His	Leu	Va]	Gln
225					230					235					240
Pro	Asp	Val	Asp	G1 y	Leu	G] u	Ser	Ser	Pro	Gly	Asp	Ala	Gln	Gln	Trp
				245					250					255	
Gly	Arg	Leu	Gln	Leu	Ser	Leu	Glu	Phe	Asp	Phe	Gly	Ser	Gln	Glu	Val
			260					265					270		
Lys	Gly	Pro	Ala	Ala	Gln	Asp	Gln	Arg	Phe	Cys	Glu	Phe	Pro	Glu	Arg
		275					280					285			
Val	Thr	Gly	Glu	Gly	Gln	Thr	Pro	Cys	Pro	G1 y	Trp	Trp	Gly	Ala	Asp
	290					295					300				
Arg	Ala	Gly	Ala	Leu	Gly										
305					310										
	0> 48														
	1> 8′														
	2> PI														
<21	3> H	omo .	sapı	ens											
/40	0> 48	00 <i>4</i>													
			A1a	116	مال	Leu	Leu	Ala	Len	len	G1 v	Ala	Met	Ser	Gly
1	OIII	Gru	MIG	5	110	Leu	LCu	MIG	10	Leu	O. y	,,,,,	me c	15	01)
	Val	Ala	Glu		Ser	Pro	Pro	Glv		Ser	Val	His	Lvs		Ser
71.511			20					25					30		
Val	Lvs	Leu			Ser	Leu	Ser	Pro	Val	He	Pro	G1 y	Phe	Pro	Gln
-	-, -	35					40					45			
lle	Val	Asn	Ser	Asn	Pro	Leu	Thr	Glu	Ala	Phe	Arg	Val	Asn	Trp	Leu
	50					55					60				
Ser	Gly	Thr	Tyr	Phe	Glu	Val	Val	Thr	Thr	Gly	Met	Glu	Gln	Leu	Asp
65					70					75					80
Phe	Glu	Thr	Gly	P.ro	Asn	lle	Phe	Asp	Leu	G1n	lle	Tyr	Val	Lys	Asp
				85					90					95	
Glu	Va]	Gly	Va1	Thr	Asp	Leu	G1n	Val	Leu	Thr	Val	Gln	Val	Thr	Asp

			100					105					110		
Val	Asn	Glu	Pro	Pro	Gln	Phe	Gln	Gly	Asn	Leu	Ala	Glu	Gly	Leu	His
		115					120					125			
Leu	Tyr	He	Val	Glu	Arg	Ala	Asn	Pro	Gly	Phe	He	Tyr	Gln	Val	G1 u
	130					135					140				
Ala	Phe	Asp	Pro	Glu	Asp	Thr	Ser	Arg	Asn	lle	Pro	Leu	Ser	Tyr	Phe
145					150					155					160
Leu	He	Ser	Pro	Pro	Lys	Ser	Phe	Arg	Met	Ser	Ala	Asn	Gly	Thr	Leu
				165					170					175	
Phe	Ser	Thr	Thr	Glu	Leu	Asp	Phe	Glu	Ala	Arg	His	Arg	Ser	Phe	His
			180					185					190		
Leu	He	Val	Glu	Val	Arg	Asp	Ser	Gly	Gly	Leu	Lys	Ala	Ser	Thr	Glu
		195					200					205			
Leu	Gln	Val	Asn	He	Val	Asn	Leu	Asn	Asp	Glu	Val	Pro	Arg	Phe	Thr
	210					215					220				
Ser	Pro	Thr	Arg	Val	Tyr	Thr	Va]	Leu	Glu	Glu	Leu	Ser	Pro	Gly	Thr
225					230					235					240
lle	Val	Ala	Asn	Ile	Thr	Ala	Glu	Asp	Pro	Asp	Asp	Glu	Gly	Phe	Pro
				245					250					255	
Ser	His	Leu	Leu	Tyr	Ser	Ile	Thr	Thr	Val	Ser	Lys	Tyr	Phe	Met	lle
			260					265					270		
Asn	Gln	Leu	Thr	Gly	Thr	He	Gln	Val	Ala	Gln	Arg	He	Asp	Arg	Asp
		275					280					285			
Ala	Gly	Glu	Leu	Arg	Gln	Asn	Pro	Thr	lle	Ser	Leu	Glu	Val	Leu	Val
	290					295					300				
Lys	Asp	Arg	Pro	Tyr	Gly	Gly	Gln	Glu	Asn	Arg	Ile	Gln	Ile	Thr	Phe
305					310					315					320
He	Val	Glu	Asp	Val	Asn	Asp	Asn	Pro	Ala	Thr	Cys	Gln	Lys	Phe	Thr
				325					330					335	
Phe	Ser	He	Met	Val	Pro	Glu	Arg	Thr	Ala	Lys	G1 y	Thr	Leu	Leu	Leu
			340					345					350		
Asp	Leu	Asn	Lys	Phe	Cys	Phe	Asp	Asp	Asp	Ser	Glu	Лlа	Pro	Asn	Asn
		355					360					365			
Arg		Asn	Phe	Thr	Met		Ser	Gly	Val	Gly		Gly	Ser	Arg	Phe
	370					375					380				
Len	Glp	Asn	Pro	Ala	G1v	Ser	G1v	lve	He	Val	Leu	11e	Glv	Asn	Len

385					390					395					400
Asp	Tyr	Glu	Asn	Pro	Ser	Asn	Leu	Ala	Ala	Gly	Asn	Lys	Tyr	Thr	Val
				405					410					415	
11e	lle	Gln	Val	Gln	Asp	Val	Ala	Pro	Pro	Tyr	Tyr	Lys	Asn	Asn	Val
			420					425					430		
Tyr	Val	Tyr	He	Leu	Thr	Ser	Pro	Glu	Asn	Glu	Phe	Pro	Leu	lle	Phe
		435					440					445			
Asp	Arg	Pro	Ser	Tyr	Val	Phe	Asp	Val	Ser	Glu	Arg	Arg	Pro	Ala	Arg
	450					455					460				
Thr	Arg	Val	Gly	Gln	Val	Arg	Ala	Thr	Asp	Lys	Asp	Leu	Pro	Gln	Ser
465					470					475					480
Ser	Leu	Leu	Tyr	Ser	He	Ser	Thr	Gly	Gly	Ala	Ser	Leu	Gln	Tyr	Pro
				485					490					495	
Asn	Val	Phe	Trp	He	Asn	Pro	Lys	Thr	G1 y	Glu	Leu	Gln	Leu	Val	Thr
			500					505					510		
Lys	Va]	Asp	Cys	Glu	Thr	Thr	Pro	He	Tyr	He	Leu	Arg	Ile	Gln	Ala
		515					520					525			
Thr	Asn	Asn	Glu	Asp	Thr	Ser	Ser	Val	Thr	Val	Thr	Val	Asn	lle	Leu
	530					535					540				
Glu	Glu	Asn	Asp	Glu	Lys	Pro	lle	Cys	Thr	Pro	Asn	Ser	Tyr	Phe	Leu
545					550					555					560
Ala	Leu	Pro	Val	Asp	Leu	Lys	Val	Gly	Thr	Asn	He	Gln	Asn	Phe	Lys
				565					570	•				575	
Leu	Thr	Cys	Thr	Asp	Leu	Asp	Ser	Ser	Pro	Arg	Ser	Phe	Arg	Tyr	Ser
			580					585					590		
He	Gly	Pro	Gly	Asn	Val	Asn	Asn	His	Phe	Thr	Phe		Pro	Asn	Ala
		595										605		_	
Gly			Val	Thr	Arg			Leu	Thr	Ser			Asp	Tyr	Ala
	610					615		_			620		_		en)
	Gly	Phe	Asp	Lys			Asp	Tyr	Lys			Val	Tyr	Val	
625					630		_			635			,	17 1	640
Asp	Asp	Asn	Leu			Asp	Lys	Lys			Glu	Ala	Leu		
				645		6			650		D		D	655	
Thr	Gly	Ihr			Leu	Ser	He			116	Pro	H1S	Pro		ınr
		an.	660		D		В	665		TI	т	C1	670 Val		Λ
110	110	The	Thr	Lhr	Pro	Arre	ピャっ	Arco	· Val	lbr	1.77	· L.In	เหลเ	1 611	Aro

Lys Asn Val Tyr Ser Pro Ser Ala Trp Tyr Val Pro Phe Val Ile Thr Leu Gly Ser Ile Leu Leu Leu Gly Leu Leu Val Tyr Leu Val Val Leu Leu Ala Lys Ala Ile His Arg His Cys Pro Cys Lys Thr Gly Lys Asn Lys Glu Pro Leu Thr Lys Lys Gly Glu Thr Lys Thr Ala Glu Arg Asp Val Val Val Glu Thr Ile Gln Met Asn Thr Ile Phe Asp Gly Glu Ala lle Asp Pro Val Thr Gly Glu Thr Tyr Glu Leu Asn Ser Lys Thr Gly Ala Arg Lys Trp Lys Asp Pro Leu Thr Gln Met Pro Lys Trp Lys Glu Ser Ser His Gln Gly Ala Ala Pro Arg Arg Val Thr Ala Gly Glu Gly Met Gly Ser Leu Arg Ser Ala Asn Trp Glu Glu Asp Glu Leu Ser Gly Lys Ala Trp Ala Glu Asp Ala Asp Leu Gly Ser Arg Asn Glu Gly Gly Lys Leu Gly Asn Pro Lys Asn Arg Asn Pro Ala Phe Met Asn Arg Ala Tyr Pro Lys Pro His Pro Gly Lys

<210> 4805

<211> 530

<212> PRT

<213> Homo sapiens

<400> 4805

Met Glu Gly Pro Leu Thr Pro Pro Pro Leu Gln Gly Gly Gly Ala Ala

1 5 10 15

Ala Val Pro Glu Pro Gly Ala Arg Gln His Pro Gly His Glu Thr Ala

Ala	Gln		Tyr	Ser	Ala	Arg		Leu	Gln	Ala	G1 y		Glu	Pro	Glu
		35					40					45			
Ser	Asp	Phe	Leu	lle	Leu	Pro	Gly	Phe	Пе	Asp	Phe	He	Ala	Asp	Glu
	50					55					60				
Val	Asp	Leu	Thr	Ser	Ala	Leu	Thr	Arg	Lys	He	Thr	Leu	Lys	Thr	Pro
65					70					75					80
Leu	Ile	Ser	Ser	Pro	Met	Asp	Thr	Val	Thr	Glu	Ala	Asp	Met	Ala	Phe
				85					90					95	
Ala	Met	Ala	Leu	Met	Gly	Gly	lle	Gly	Phe	lle	His	His	Asn	Cys	Thr
			100					105					110		
Pro	Glu	Phe	Gln	Ala	Asn	Glu	Val	Arg	Lys	Val	Lys	Lys	Phe	Glu	Gln
		115					120					125			
Gly	Phe	He	Thr	Asp	Pro	Val	Val	Leu	Ser	Pro	Ser	His	Thr	Val	Gly
	130					135					140				
Asp	Val	Leu	Glu	Ala	Lys	Met	Arg	His	Gly	Phe	Ser	Gly	lle	Pro	lle
145					150					155					160
Thr	Glu	Thr	Gly	Thr	Met	Gly	Ser	Lys	Leu	Val	Gly	Ile	Val	Thr	Ser
				165					170					175	
Arg	Asp	Ile	Asp	Phe	Leu	Ala	Glu	Lys	Asp	His	Thr	Thr	Leu	Leu	Ser
J	•		180					185					190		
Glu	Val	Met		Pro	Arg	He	Glu	Leu	Val	Val	Ala	Pro	Ala	Gly	Val
		195					200					205			
Thr	Leu		Glu	Ala	Asn	Glu		Leu	Gln	Arg	Ser		Lvs	Glv	Lvs
	210					215					220	•	•		
Leu		He	Val	Asn	Asp			Glu	l.eu	Val		lle	He	Ala	Arg
225					•	•	_								240
		Leu	Lvs	Lvs	Asn	Arg	Asp	Tyr	Pro	l.eu	Ala	Ser	Lvs	Asp	Ser
	пор	Вса	25,0	245		6		-,-	250					255	
Gln	lve	Gln	Len		Cys	Glv	Ala	Ala			Thr	Arø	Glu		Asp
OIN	Lys	0111	260	LCu	0,5	0.1 9	7110	265		0.1)		6	270		
Lve	Tur	Ara		Aen	Leu	ا ما	Thr		Ala	Glu	Val	Aen]]_	Val
Lys	1 y 1	275	Leu	nsp	Leu	126.17	280	9111	1310	O į į	• 61	285			. (1)
1	Λ		Can	C1 _m	Gly	Acr		Vo.1	Ту	616	110			Va1	Hic
rea	nsp	ser	261	OIII	OIA	ASII	Sel	val	1 y 1	0111	116	MIG	MC L	· a I	111.5

```
Tyr Ile Lys Gln Lys Tyr Pro His Leu Gln Val Ile Gly Gly Asn Val
305
                    310
                                        315
Val Thr Ala Ala Gln Ala Lys Asn Leu lle Asp Ala Gly Val Asp Gly
                325
                                    330
Leu Arg Val Gly Met Gly Cys Gly Ser Ile Cys Ile Thr Gln Glu Val
                                345
Met Ala Cys Gly Arg Pro Gln Gly Thr Ala Val Tyr Lys Val Ala Glu
                            360
                                                 365
Tyr Ala Arg Arg Phe Gly Val Pro Ile Ile Ala Asp Gly Gly Ile Gln
                        375
                                             380
Thr Val Gly His Val Val Lys Ala Leu Ala Leu Gly Ala Ser Thr Val
                    390
                                        395
Met Met Gly Ser Leu Leu Ala Ala Thr Thr Glu Ala Pro Gly Glu Tyr
                405
                                    410
                                                         415
Phe Phe Ser Asp Gly Val Arg Leu Lys Lys Tyr Arg Gly Met Gly Ser
            420
                                425
Leu Asp Ala Met Glu Lys Ser Ser Ser Ser Gln Lys Arg Tyr Phe Ser
                            440
Glu Gly Asp Lys Val Lys Ile Ala Gln Gly Val Ser Gly Ser Ile Gln
    450
                        455
Asp Lys Gly Ser Ile Gln Lys Phe Val Pro Tyr Leu Ile Ala Gly Ile
                                        475
                    470
Gln His Gly Cys Gln Asp 11e Gly Ala Arg Ser Leu Ser Val Leu Arg
                                    490
                485
                                                         495
Ser Met Met Tyr Ser Gly Glu Leu Lys Phe Glu Lys Arg Thr Met Ser
            500
                                505
                                                     510
Ala Gln Ile Glu Gly Gly Val His Gly Leu His Ser Tyr Glu Lys Arg
                            520
                                                 525
Leu Tyr
    530
```

<210> 4806

<211> 134

<212> PRT

<213> Homo sapiens

<400> 4806 Met Leu Arg Asp Phe Val Thr Thr Arg Pro Ala Leu Gln Glu Ala Leu 1 5 10 15 Lys Glu Ala Leu Asn Met Glu Arg Lys Asn Arg Tyr Gln Pro Leu Gln 25 Lys Pro Tyr Gln Asn Met Lys Thr Asn Glu Ala Ile Lys Lys Leu His 40 45 Gln Leu Met Cys Lys Ile Thr Arg Ser Asn Ser His Ile Thr Ile Leu 50 Thr Leu Ser Val Asn Trp Leu Asn Ala Pro Ile Lys Lys His Arg Leu 70 75 Ala Asn Trp lle Lys Ser Gln Asp Pro Ser Val Cys Cys Ile Gln Glu 85 90 95 Thr His Leu Met Cys Lys Asp Thr His Arg Pro Lys Ile Lys Gly Trp 100 105 Arg Asn Ile Tyr Gln Ala Asn Gly Lys Pro Lys Lys Lys Ser Gly 120 115 125 Gly Cys Asn Pro Ser Leu 130 <210> 4807 <211> 505 <212> PRT <213> Homo sapiens <400> 4807 Met Ala Gly Ala Gln Thr Leu Leu Thr Phe Arg Asp Val Ala lle Glu 10 Phe Ser Leu Glu Glu Trp Lys Cys Leu Asp Leu Ala Gln Gln Asn Leu 25 Tyr Arg Asp Val Met Leu Glu Asn Tyr Arg Asn Leu Phe Ser Val Gly

40

55

Leu Thr Val Cys Lys Pro Gly Leu 11e Thr Cys Leu Glu Gln Arg Lys

45

60

35

Glu	Pro	Trp	Asn	Val	Lys	Arg	Gln	Glu	Ala	Ala	Asp	Gly	His	Pro	Ala
65					70					75					80
Met	Ser	Ser	His	Phe 85	Thr	Gln	Asp	Leu	Leu 90	Pro	Glu	G1n	G1y	11e 95	G1n
Asp	Ala	Phe	Pro 100	Lys	Arg	lle	Leu	Arg 105	Gly	Tyr	Gly	Asn	Cys 110	Gly	Leu
Asp	Asn	Leu 115	Tyr	Leu	Arg	Lys	Asp 120	Trp	Glu	Ser	Leu	Аsp 125	Glu	Cys	Lys
Leu	Gln 130	Lys	Asp	Tyr	Asn	Gly 135	Leu	Asn	Gln	Cys	Ser 140	Ser	Thr	Thr	His
Ser 145	Lys	Ile	Phe	Gln	Tyr 150	Asn	Lys	Tyr	Val	Lys 155	Ile	Phe	Asp	Asn	Phe 160
Ser	Asn	Leu	His	Arg 165	Arg	Asn	lle	Ser	Asn 170	Thr	Gly	Glu	Lys	Pro 175	Phe
Lys	Cys	G1n	Glu 180	Cys	Gly	Lys	Ser	Phe 185	Gln	Met	Leu	Ser	Phe 190	Leu	Thr
Glu	His	G1n 195	Lys	lle	His	Thr	Gly 200	Lys	Lys	Phe	Gln	Lys 205	Cys	Gly	Glu
Cys	Gly 210	Lys	Thr	Phe	Ile	Gln 215	Cys	Ser	His	Phe	Thr 220	Glu	Pro	Glu	Asn
Ile 225	Asp	Thr	Gly	Glu	Lys 230	Pro	Tyr	Lys	Cys	Gln 235	Glu	Cys	Asn	Asn	Val 240
lle	Lys	Thr	Cys	Ser 245	Val	Leu	Thr	Lys	Asn 250	Arg	lle	Tyr	Ala	Gly 255	Gly
Glu	His	Tyr	Arg 260	Cys	Glu	Glu	Phe	Gly 265	Lys	Va]	Phe	Asn	Gln 270	Cys	Ser
His	Leu	Thr 275	Glu	His	Glu	His	Gly 280	Thr	Glu	Glu	Lys	Pro 285	Phe	Lys	Cys
Glu	Glu 290	Cys	Asp	Ser	lle	Phe 295	Lys	Trp	Phe	Ser	Asp 300	Leu	Thr	Lys	His
Lys 305	Arg	lle	His	Thr	Gly 310	Glu	Lys	Pro	Tyr	Lys 315	Cys	Asp	Glu	Cys	G1y 320
Lys	Ala	Tyr	Thr	Gln 325	Ser	Ser	llis	Leu	Ser 330	Glu	His	Arg	Arg	11e 335	His
Thr	Gly	Glu	Lys	Pro	Tyr	Gln	Cys	Glu	Glu	Cys	Gly	Lys	Val	Phe	Arg

Thr Cys Ser Ser Leu Ser Asn His Lys Arg Thr His Ser Glu Glu Lys Pro Tyr Thr Cys Glu Glu Cys Gly Asn Ile Phe Lys Gln Leu Ser Asp Leu Thr Lys His Lys Lys Thr His Thr Gly Glu Lys Pro Tyr Lys Cys Asp Glu Cys Gly Lys Asn Phe Thr Gln Ser Ser Asn Leu Ile Val His Lys Arg Ile His Thr Gly Glu Lys Pro Tyr Lys Cys Glu Glu Cys Gly Arg Val Phe Met Trp Phe Ser Asp Ile Thr Lys His Lys Lys Thr His Thr Gly Glu Lys Pro Tyr Lys Cys Asp Glu Cys Gly Lys Asn Phe Thr Gln Ser Ser Asn Leu Ile Val His Lys Arg lle His Thr Gly Glu Lys Pro Tyr Lys Cys Glu Lys Cys Gly Lys Ala Phe Thr Gln Phe Ser His Leu Thr Val His Glu Ser Ile His Thr

<210> 4808

<211> 106

<212> PRT

<213> Homo sapiens

<400> 4808

 Met Val Ala Pro Arg Lys Ile Arg Trp Cys Val Glu Glu Leu Pro Ser

 1
 5
 10
 15

 His Cys Ser Leu His Ala Phe Leu Ser Ala Arg Cys Glu Gln Gly Arg
 20
 25
 30

 Pro Gly Thr Ala Asp Gly Leu His Leu Ser Asp Trp Thr Ala Thr Cys
 45

 His Thr Phe Ile Glu Leu Ser Ile Ser Thr Ala Ala Val Leu Gly Val
 50
 55

 Leu Leu Gly Ser Arg
 Arg Asn Lys Lys Phe Arg Glu Ala Lys Val Leu Gln

 65
 70
 75
 80

 Gly Cys Arg Gln Val Ile Glu Pro Ser Phe Ile Glu His Leu Leu Ser

 85
 90
 95

 Ala Lys Tyr Ser Tyr Leu Arg Tyr Leu Val

 100
 105

<210> 4809

<211> 410

<212> PRT

<213> Homo sapiens

<400> 4809

Met Phe Val Trp Asn Gln Gly Ser Lys Tyr Ser Gly Ser Glu Val Leu 1 5 10 15

Trp Cys Pro Leu His Trp Asn Ser Glu Glu Leu Ile Val Phe Trp Ser 20 25 30

Cys Val Leu Gln Val Leu Asp Trp His Arg Leu Ile Pro Leu Thr Trp
35 40 45

Ala Cys Met Ala Arg Gln Thr Pro His Leu Gly Glu Gln Arg Arg Thr
50 55 60

Thr Ala Ser Leu Leu Arg Lys Leu Thr Thr Ala Ser Asn Gly Gly Val 65 70 75 80

lle Glu Glu Leu Ser Cys Val Arg Ser Asn Asn Tyr Val Gln Glu Pro 85 90 95

Glu Cys Arg Arg Asn Leu Val Gln Cys Leu Leu Glu Lys Gln Gly Thr 100 105 110

Pro Val Val Gln Gly Ser Leu Glu Leu Glu Arg Val Met Ser Ser Leu 115 120 125

Leu Asp Met Gly Phe Ser Asn Ala His Ile Asn Glu Leu Leu Ser Val 130 135 140

Arg Arg Gly Ala Ser Leu Gln Gln Leu Leu Asp Ile 1le Ser Glu Phe 145 150 155 160

lle Leu Leu Gly Leu Asn Pro Glu Pro Val Cys Val Val Leu Lys Lys 165 170 175

Ser	Pro	Gln	Leu	Leu	Lys	Leu	Pro	lle	Met	Gln	Met	Arg	Lys	Arg	Ser
			180					185					190		
Ser	Tyr	Leu	G1n	Lys	Leu	Gly	Leu	Gly	Glu	Gly	Lys	Leu	Lys	Arg	Val
		195					200					205			
Leu	Tyr	Cys	Cys	Pro	Glu	lle	Phe	Thr	Met	Arg	Gln	Gln	Asp	He	Asn
	210					215					220				
Asp	Thr	Val	Arg	Leu	Leu	Lys	Glu	Lys	Cys	Leu	Phe	Thr	Val	Gln	Gln
225					230					235					240
Val	Thr	Lys	Ile	Leu	His	Ser	Cys	Pro	Ser	Val	Leu	Arg	Glu	Asp	Leu
				245					250					255	
Gly	Gln	Leu	Glu	Tyr	Lys	Phe	Gln	Tyr	Ala	Tyr	Phe	Arg	Met	Gly	Ile
			260					265					270		
Lys	His	Pro	Asp	lle	Val	Lys	Ser	Glu	Tyr	Leu	G1n	Tyr	Ser	Leu	Thr
		275					280				-	285			
Lys	He	Lys	Gln	Arg	His	He	Tyr	Leu	Glu	Arg	Leu	Gly	Arg	Tyr	Gln
	290					295					300				
Thr	Pro	Asp	Lys	Lys	Gly	Gln	Thr	Gln	He	Pro	Asn	Pro	Leu	Leu	Lys
305					310					315					320
Asp	Ile	Leu	Arg	Val	Ser	Glu	Ala	Glu	Phe	Leu	Ala	Arg	Thr	Ala	Cys
				325					330					335	
Thr	Ser	Val	Glu	Glu	Phe	Gln	Val	Phe	Lys	Lys	Leu	Leu	Ala	Arg	Glu
			340					345					350		
G1u	Glu	Glu	Ser	Glu	Ser	Ser	Thr	Ser	Asp	Asp	Lys	Arg	Ala	Ser	Leu
		355					360					365			
Asp	Glu	Asp	Glu	Asp	Asp	Asp	Asp	Glu	Glu	Asp	Asn	Asp	Glu	Asp	Asp
	370					375					380				
Asn	Asp	Glu	Asp	Asp	Asp	Asp	Glu	Asp	Asp	Asp	Glu	Ala	Glu	Asp	Asn
385					390					395					400
Asp	Glu	Asp	Glu	Asp	Asp	Asp	Glu	G] u	Glu						
				405					410						

<211> 340

<212> PRT

<213> Homo sapiens

<400)> 48	310													
Met	Ala	Met	Gln	Met	Gln	Leu	Glu	Ala	Asn	Ala	Asp	Thr	Ser	Val	Glu
1				5					10					15	
Glu	Glu	Ser	Phe	Gly	Pro	Gln	Pro	lle	Ser	Arg	Leu	G]u	Gln	Cys	Gly
			20					25					30		
lle	Asn	Ala	Asn	Asp	Val	Lys	Lys	Leu	Glu	Glu	Ala	Gly	Phe	His	Thr
		35					40					45			
Val	Glu	Ala	Val	Ala	Tyr	Ala	Pro	Lys	Lys	Glu	Leu	He	Asn	Ile	Lys
	50					55					60				
Gly	Ile	Ser	Glu	Ala	Lys	Ala	Asp	Lys	Ile	Leu	Thr	Glu	Ser	Arg	Ser
65					70					75					80
Val	Ala	Arg	Leu	Glu	Cys	Asn	Ser	Val	He	Leu	Val	Tyr	Cys	Thr	Leu
				85					90					95	
Arg	Leu	Ser	Gly	Ser	Ser	Asp	Ser	Pro	Ala	Ser	Ala	Ser	Arg	Val	Val
			100					105					110		
Gly	Thr	Thr	Gly	Gly	Ile	Glu	Thr	Gly	Ser	Ile	Thr	Glu	Met	Phe	Gly
		115					120					125			
Glu	Phe	Arg	Thr	Gly	Lys	Thr	G]n	Ile	Cys	His	Thr	Leu	Ala	Val	Thr
	130					135					140				
Cys	Gln	Leu	Pro	He	Asp	Arg	Gly	Gly	Gly		Gly	Lys	Ala	Met	
145					150					155					160
He	Asp	Thr	Glu		Thr	Phe	Arg	Pro		Arg	Leu	Leu	Ala		Ala
		_		165			~		170				., .	175	m
Glu	Arg	Tyr		Leu	Ser	GI y	Ser		Val	Leu	Asp	Asn	Val	Ala	lyr
		. 1	180		T)			185	TI	61	,		190	C1	A 1 .
Ala	Arg		Phe	Asn	Ihr	Asp		GIn	ınr	GIn	Leu		Tyr	61n	Ala
c	4.1	195	14 .	W = 1	C1	C	200	Т	A 1	1	Lan	205		Aan	Cox
Ser			Met	vai	61u			lyr	ATA	Leu			Va]	ASP	Sei
	210					215					220				
110	Tha	Λla	Lou	Tun	Ara	Thr	Aco	Tur	Sor	GLv	Ara	Glv	Glu	Lou	Sar
		Ма	Leu	1 7 1	230	1111	nsp	1 9 1	361	235		ОГУ	Olu	Leu	240
225		Gla	Mot	Hic		Ala	Ara	Pho	الما			الم [Leu	Arø	
nia	un g	OIII	me t	245	Leu	111a	m g	1116	250		MC L	200	. "Cu	255	Doc
41.	۸	C1	Dl. a	C1	V a 1	41.	Va 1	Vo.1		Tha	Aan	C1n	Vol		۸1،

			260					265					270		
Gln	Val	Asp 275	Gly	Ala	Ala	Met	Phe 280	Ala	Ala	Asp	Pro	Lys 285	Lys	Pro	Ile
Gly		Asn	lle	Ile	Ala	His	Ala	Ser	Thr	Thr	Arg	Leu	Tyr	Leu	Arg
lvs	290 Glv	Arø	Glv	Glu	Thr	295 Arg	He	Cvs	Lvs	He		Asp	Ser	Pro	Cvs
305	01)	111 6	01)	014	310	*** 8	110	Cyc	2,0	315	.,.				320
	Pro	Glu	Ala	Glu		Met	Phe	Ala	He		Ala	Asp	Gly	Val	
				325					330					335	
Asp	Ala	Lys	Asp												
			340												
)> 48														
	1> 69														
	2> PI														
₹21.	32 H	omo :	sapi	ens											
<400)> 48	811													
			Arg	Gly	Leu	Ala	Asp	Ser	Gly	Gln	Gly	Ser	Phe	Thr	Gly
			Arg	Gly 5	Leu	Ala	Asp	Ser	Gly 10	Gln	Gly	Ser	Phe	Thr 15	Gly
Met 1	Glu	Thr		5		Ala			10					15	
Met 1	Glu	Thr		5					10					15	
Met 1 Gln	Glu Gly	Thr	Ala 20	5 Arg	Phe		Arg	11e 25	10 Gln	Lys	Lys	Ser	Gln 30	15 Pro	Glu
Met 1 Gln Lys	Glu Gly Val	Thr lle Val	Ala 20 Arg	5 Arg Ala	Phe Ala	Gly Ser	Arg Arg 40	lle 25 Gly	10 Gln Arg	Lys Pro	Lys Leu	Ser Ile 45	Gln 30 Gly	15 Pro Trp	Glu Thr
Met 1 Gln Lys	Glu Gly Val	Thr lle Val	Ala 20 Arg	5 Arg Ala	Phe Ala	Gly	Arg Arg 40	lle 25 Gly	10 Gln Arg	Lys Pro	Lys Leu	Ser Ile 45	Gln 30 Gly	15 Pro Trp	Glu Thr
Met 1 Gln Lys	Glu Gly Val	Thr lle Val	Ala 20 Arg	5 Arg Ala	Phe Ala	Gly Ser	Arg Arg 40 Gly	lle 25 Gly	10 Gln Arg	Lys Pro	Lys Leu	Ser Ile 45 Met	Gln 30 Gly	15 Pro Trp	Glu Thr
Met 1 Gln Lys Gln	Glu Gly Val Trp 50	Thr 11e Val 35 Cys	Ala 20 Arg Ala	5 Arg Ala Glu	Phe Ala Asp	Gly Ser Gly 55	Arg Arg 40 Gly	lle 25 Gly Asp	10 Gln Arg Glu	Lys Pro Ser	Lys Leu Glu 60	Ser Ile 45 Met	Gln 30 Gly Ala	15 Pro Trp Leu	Glu Thr
Met 1 Gln Lys Gln	Glu Gly Val Trp 50	Thr 11e Val 35 Cys	Ala 20 Arg Ala	5 Arg Ala Glu	Phe Ala Asp	Gly Ser	Arg Arg 40 Gly	lle 25 Gly Asp	10 Gln Arg Glu	Lys Pro Ser	Lys Leu Glu 60	Ser Ile 45 Met	Gln 30 Gly Ala	15 Pro Trp Leu	Glu Thr Ala
Met 1 Gln Lys Gln Gly 65	Glu Gly Val Trp 50 Ser	Thr lle Val 35 Cys	Ala 20 Arg Ala Gly	5 Arg Ala Glu Cys	Phe Ala Asp Ser 70	Gly Ser Gly 55 Ser	Arg Arg 40 Gly	lle 25 Gly Asp	10 Gln Arg Glu	Lys Pro Ser Gly 75	Lys Leu Glu 60 Arg	Ser Ile 45 Met	Gln 30 Gly Ala	15 Pro Trp Leu	Glu Thr Ala Leu 80
Met 1 Gln Lys Gln Gly 65	Glu Gly Val Trp 50 Ser	Thr lle Val 35 Cys	Ala 20 Arg Ala Gly	5 Arg Ala Glu Cys	Phe Ala Asp Ser 70	Gly Ser Gly 55	Arg Arg 40 Gly	lle 25 Gly Asp	10 Gln Arg Glu	Lys Pro Ser Gly 75	Lys Leu Glu 60 Arg	Ser Ile 45 Met	Gln 30 Gly Ala	15 Pro Trp Leu	Glu Thr Ala Leu 80
Met 1 Gln Lys Gln Gly 65 Ile	Glu Gly Val Trp 50 Ser	Thr 11e Val 35 Cys Pro	Ala 20 Arg Ala Gly Leu	5 Arg Ala Glu Cys Arg 85	Phe Ala Asp Ser 70 Arg	Gly Ser Gly 55 Ser	Arg Arg 40 Gly Gly	lle 25 Gly Asp Pro	10 Gln Arg Glu Gln Arg 90	Lys Pro Ser Gly 75 His	Lys Leu Glu 60 Arg	Ser Ile 45 Met	Gln 30 Gly Ala Ser	15 Pro Trp Leu Arg Gln 95	Glu Thr Ala Leu 80
Met 1 Gln Lys Gln Gly 65 Ile	Glu Gly Val Trp 50 Ser	Thr 11e Val 35 Cys Pro	Ala 20 Arg Ala Gly Leu	5 Arg Ala Glu Cys Arg 85	Phe Ala Asp Ser 70 Arg	Gly Ser Gly 55 Ser	Arg Arg 40 Gly Gly	lle 25 Gly Asp Pro	10 Gln Arg Glu Gln Arg 90	Lys Pro Ser Gly 75 His	Lys Leu Glu 60 Arg	Ser Ile 45 Met	Gln 30 Gly Ala Ser	15 Pro Trp Leu Arg Gln 95	Glu Thr Ala Leu 80
Met 1 Gln Lys Gln Gly 65 Ile Gln	Glu Gly Val Trp 50 Ser Phe Gly	Thr 11e Val 35 Cys Pro Leu Pro	Ala 20 Arg Ala Gly Leu Asp 100	5 Arg Ala Glu Cys Arg 85 Ser	Phe Ala Asp Ser 70 Arg	Gly Ser Gly 55 Ser	Arg 40 Gly Ala Asp	Ile 25 Gly Asp Pro Ala Arg 105	10 Gln Arg Glu Gln Arg 90 Phe	Lys Pro Ser Gly 75 His	Lys Leu Glu 60 Arg Val	Ser Ile 45 Met Leu His	Gln 30 Gly Ala Ser His Glu 110	15 Pro Trp Leu Arg Gln 95 Leu	Glu Thr Ala Leu 80 Asp

Glu		Ala	Asp	Arg	Gly		Ser	Ala	Trp	Pro	Leu 140	Ala	Lys	Cys	Asn
Thm	130	The	Con	Aan	Aan	135	Glu	G1 ₁₁	C1	Lys		Thr	Lve	Lve	lve
	ASII	HIL	ser	ASII		1111	Giu	Gru	ULU	155	Lys	1 1 1 1 1	Lys	Lys	160
145	A1.	11.	Val	Val	150	Dro	San	San	Aan		Tur	Tur	Ara	Trn	
ASP	мта	116	val		изр	110	261	261	170	Leu	1 9 1	1 9 1	Mg	175	Leu
Tha	A10	11.	110	165	Dno	Val	Dho	Hic		Trp	Tyr	Lou	Lou		Cvc
IIII	нта	116	180	Leu	110	Val	rne	185	ASII	чи	1 9 1	Leu	190	110	Cys
Ara	110	Cvc		Acn	Glu	Lou	Gln		Glu	Tyr	Lau	Mot		Trn	Leu
Arg	на	195	rne	wsb	Giu	Leu	200	261	Ulu	1 9 1	Leu	205	Leu	пр	Leu
Vol	Lou		Tur	Sor	Δla	Acn		Lou	Tur	Val	Leu		Val	Len	Val
vai	210	пър	1 9 1	Jei	MIG	215	101	LCu	1 9 1	,41	220	пор	, 41	Lea	, 41
Ara		Aro	Thr	Glv	Phe		Glu	Gln	Glv	Leu		Val	Ser	Asp	Thr
225	MIG	m g	1111	01,	230	1500	Olu	0111	01,	235			00.	пор	240
	Arg	Leu	Trp	Gln		Tvr	Lvs	Thr	Thr	Thr	Gln	Phe	Lvs	Leu	
71071	6	,,,,,		245	.,	- , -	, -	• • • • • • • • • • • • • • • • • • • •	250				_,	255	•
Val	Leu	Ser	Leu		Pro	Thr	Asp	Leu		Tyr	Leu	Lys	Val	Gly	Thr
			260				•	265					270		
Asn	Tyr	Pro	Glu	Val	Arg	Phe	Asn	Arg	Leu	Leu	Lys	Phe	Ser	Arg	Leu
		275					280					285			
Phe	Glu	Phe	Phe	Asp	Arg	Thr	Glu	Thr	Arg	Thr	Asn	Tyr	Pro	Asn	Met
	290					295					300				
Phe	Arg	He	Gly	Asn	Leu	Val	Leu	Tyr	He	Leu	lle	He	lle	His	Trp
305					310					315					320
Asn	Ala	Cys	He	Tyr	Phe	Ala	lle	Ser	Lys	Phe	He	Gly	Phe	G1 y	Thr
				325					330					335	
Asp	Ser	Trp	Val	Tyr	Pro	Asn	He	Ser	He	Pro	Glu	His	Gly	Arg	Leu
			340			•		345					350		
Ser	Arg	Lys	Tyr	lle	Tyr	Ser	Leu	Tyr	Trp	Ser	Thr	Leu	Thr	Leu	Thr
		355					360					365			
Thr	lle	Gly	Glu	Thr	Pro	Pro	Pro	Val	Lys	Asp	Glu	Glu	Tyr	Leu	Phe
	370	•				375					380				
Val	Va]	Val	Asp	Phe	Leu	Val	Gly	Val	Leu	He	Phe	Ala	Thr	He	Val
385					390					395					400
Gly	Asn	Val	Gly	Ser	Met	He	Ser	Asn	Met	Asn	Ala	Ser	Arg	Ala	Glu
				405					410					415	

Phe	Gln	Ala	Lys	lle	Asp	Ser	lle	Lys	Gln	Tyr	Met	Gln	Phe	Arg	Lys
			420					425					430		
Val	Thr	Lys	Asp	Leu	Glu	Thr	Arg	Val	lle	Arg	Trp	Phe	Asp	Tyr	Leu
		435					440					445			
Trp	Ala	Asn	Lys	Lys	Thr	Val	Asp	Glu	Lys	Glu	Val	Leu	Lys	Ser	Leu
	450					455					460				
Pro	Asp	Lys	Leu	Lys	Ala	Glu	lle	Ala	11e	Asn	Val	His	Leu	Asp	Thr
465					470					475					480
Leu	Lys	Lys	Val	Arg	Ile	Phe	Gln	Asp	Cys	Glu	Ala	Gly	Leu	Leu	Val
				485					490					495	
Glu	Leu	Val	Leu	Lys	Leu	Arg	Pro	Thr	Val	Phe	Ser	Pro	Gly	Asp	Tyr
			500					505					510		
He	Cys	Lys	Lys	Gly	Asp	He	Gly	Lys	Glu	Met	Tyr	He	He	Asn	Glu
		515					520					525			
Gly	Lys	Leu	Ala	Val	Val	Ala	Asp	Asp	Gly	Val	Thr	Gln	Phe	Val	Val
	530					535					540				
Leu	Ser	Asp	Gly	Ser	Tyr	Phe	Gly	Glu	Ile	Ser	lle	Leu	Asn	lle	Lys
545					550					555					560
Gly	Ser	Lys	Ser	Gly	Asn	Arg	Arg	Thr	Ala	Asn	lle	Arg	Ser	Ile	Gly
				565					570					575	
Tyr	Ser	Asp	Leu	Phe	Cys	Leu	Ser		Asp	Asp _.	Leu	Met		Ala	Leu
			580					585					590		
Thr	Glu		Pro	Glu	Ala	Lys		Ala	Leu	Glu	Glu		Gly	Arg	Gln
		595					600					605			
He		Met	Lys	Asp	Asn	Leu	He	Asp	Glu	Glu		Ala	Arg	Ala	Gly
	610					615					620				
	Asp	Pro	Lys	Asp		Glu	Glu	Lys	Va]		Gln	Leu	Gly	Ser	
625					630					635				_	640
Leu	Asp	Thr	Leu		Thr	Arg	Phe	Ala		Leu	Leu	Ala	Glu		Asn
	m.	0.1		645			6.1		650		0.1		0.1	655	<i>a</i> 1
Ma	Thr	GIn		Lys	Met	Lys	GIn	_	Leu	Ser	GIn	Leu		Ser	GIn
		0.1	660	0.1				665			61	0.1	670	Б	0.1
Val	Lys	-	Gly	Gly	Asp	Lys		Leu	Ala	Asp	Gly		Val	Pro	Gly
	A 3,	675	,	TI	C.I	4	680	C.	61			685			
Asp		Ihr	Lys	lhr	6Ju	Asp	Lys	GIn	GIn						
	690					695									

```
<211> 544
<212> PRT
<213> Homo sapiens
<400> 4812
Met Asp Gly Leu Asp Asn Arg Gly Glu Ile Val Val Ile Gly Ala Thr
                  5
                                      10
Asn Arg Leu Asp Ser Ile Asp Pro Ala Leu Arg Arg Pro Gly Arg Phe
                                 25
Asp Arg Glu Phe Leu Phe Asn Leu Pro Asp Gln Lys Ala Arg Lys His
         35
                             40
                                                  45
lle Leu Gln lle His Thr Arg Asp Trp Asn Pro Lys Leu Ser Asp Ala
                         55
Phe Leu Gly Glu Leu Ala Glu Lys Cys Val Gly Tyr Cys Gly Ala Asp
                     70
                                          75
Ile Lys Ala Leu Cys Thr Glu Ala Ala Leu Ile Ala Leu Arg Arg Arg
                                      90
                                                          95
Tyr Pro Gln Ile Tyr Ala Ser Ser His Lys Leu Gln Leu Asp Val Ser
                                105
Ser Ile Val Leu Ser Ala Gln Asp Phe Tyr His Ala Met Gln Asn Ile
                                                 125
        115
                            120
Val Pro Ala Ser Gln Arg Ala Val Met Ser Ser Gly His Ala Leu Ser
    130
                        135
Pro lle lle Arg Pro Leu Leu Glu Arg Ser Phe Asn Asn Ile Leu Ala
                    150
                                         155
Val Leu Gln Lys Val Phe Pro His Ala Glu Ile Ser Gln Ser Asp Lys
                                     170
                                                         175
                165
Lys Glu Asp lle Glu Thr Leu lle Leu Glu Asp Ser Glu Asp Glu Asn
                                 185
Ala Leu Ser 11e Phe Glu Thr Asn Cys His Ser Gly Ser Pro Lys Lys
                            200
                                                 205
        195
Gln Ser Ser Ser Ala Ala Ile His Lys Pro Tyr Leu His Phe Thr Met
                        215
                                             220
    210
```

Ser	Pro	Tyr	His	Gln	Pro	Thr	Ser	Tyr	Arg	Pro	Arg	Leu	Leu	Leu	Ser
225					230					235					240
Gly	Glu	Arg	Gly	Ser	Gly	Gln	Thr	Ser	His	Leu	Ala	Pro	Ala	Leu	Leu
				245					250					255	
His	Thr	Leu	Glu	Arg	Phe	Ser	Val	His	Arg	Leu	Asp	Leu	Pro	Ala	Leu
			260					265					270		
Tyr	Ser	Val	Ser	Ala	Lys	Thr	Pro	Glu	Glu	Ser	Cys	Ala	Gln	He	Phe
		275					280					285			
Arg	Glu	Ala	Arg	Arg	Thr	Val	Pro	Ser	He	Val	Tyr	Met	Pro	His	Ile
	290					295					300				
Gly	Asp	Trp	Trp	Glu	Ala	Val	Ser	Glu	Thr	Val	Arg	Ala	Thr	Phe	
305					310					315					320
Thr	Leu	Leu	Gln		He	Pro	Ser	Phe		Pro	He	Phe	Leu		Ser
				325					330					335	
Thr	Ser	Glu		Met	Tyr	Ser	Glu	Leu	Pro	Glu	Glu	Val		Cys	He
			340					345					350		
Phe	Arg		Gln	Tyr	Glu	Glu		Leu	Tyr	He	Gln		Pro	He	Glu
		355					360					365			
Glu		Arg	Arg	Lys	Phe		GIn	Glu	Leu	lie		Asn	GIn	Ala	Ser
	370	Б	D			375		4.7	. 1		380	A 7		C1	17 1
	Ala	Pro	Pro	Arg		Lys	HIS	Ala	Ala		tys	Ala	Met	Glu	
385	D		. 1		390	C	15	D		395	,	C	C1 -	C	400
Leu	Pro	Leu	Ala		Pro	Ser	Pro	Pro		GIN	Leu	Ser	GIU		GIU
	C	A	Maa	405	Α	C1	C1	C1	410	ть	1	۸	C1	415	۸
LŅS	26L	Arg	ме t 420	GIU	ASP	GIN	GIU	Glu 425	ASI	Inr	Leu	Arg		Leu	Arg
Lau	Dho	Lou		Acn	Vol	The	Lvc	Arg	Lou	Alo	Thr	Acn	430	Ara	Pho
ı.eu	rne	435	AIG	АЅР	vai	1111	440	Mg	Leu	nia	1111	445	Lys	AI g	The
Acn	Ha		Sor	Lve	Pro	Val		He	Glu	Glu	Val		Asn	Tyr	Len
поп	450	TITC	561	12 y 3	110	455	изр	110	Olu	014	460	501	пор	1) 1	Leu
Glu		He	lvs	Glu	Pro		Asn	Leu	Ser	Thr		He	Thr	Lvs	He
465		110	129.0	0,10	470		, icp	Lea	001	475	, (1)	110	,	2,0	480
	lvs	His	Asn	Tvr		Thr	Ala	Lys	Asn		Len	Lvs	Asp	lle	
	٠., ٥			485	.,			2,0	490		200	٠, ٠, ٠		495	
Leu	He	Cvs	Ser		Λla	Leu	G1u	Tyr		Pro	Asp	Lvs	Asp		Glv
			500				-	505		_ •			510		-

Asp Lys Ile 1le Arg His Arg Ala Cys Thr Leu Lys Asp Thr Ala His 515 520 525

Ala 1le Ile Ala Ala Glu Leu Asp Pro Glu Phe Asn Lys Leu Cys Glu 530 535 540

<210> 4813

<211> 334

<212> PRT

<213> Homo sapiens

<400> 4813

Met Val Gly Gln Gly Cys Ser Leu Glu Pro Arg Leu Asp Leu Arg Pro

1 5 10 15

Leu Gly Asn Pro Arg Ser Lys Gln Ser Arg His Gln Ser Trp Val Trp
20 25 30

Glu Arg Gln Glu Gly Glu Arg Asn Gly Gly Ser Lys Glu Gly Lys Glu
35 40 45

Gly Gly Arg Gly Gly Ser Lys Thr Val Ile Pro Ile Pro Ile Ser Asp 50 55 60

Leu Glu Leu Ala Ser Thr Pro Val His Pro Cys Arg Gly Gly Pro Ser
65 70 75 80

Pro Gln Leu Leu Pro Arg Glu Tyr Gly Gly Gly Val Gly Ile Leu Trp 85 90 95

Leu Ser Ser Leu Pro Leu Leu Cys Arg Leu Met Leu Gly Phe Met Gly
100 105 110

Val Thr Ala Leu Leu Ser Met Trp Ile Ser Asn Thr Ala Thr Thr Ala 115 120 125

Met Met Val Pro lle Val Glu Ala Ile Leu Gln Gln Met Glu Ala Thr 130 135 140

Ser Ala Ala Thr Glu Ala Gly Leu Glu Leu Val Asp Lys Gly Lys Ala 145 150 155 160

Lys Glu Leu Pro Gly Ser Gln Val Ile Phe Glu Gly Pro Thr Leu Gly
165 170 175

Gln Gln Glu Asp Gln Glu Arg Lys Arg Leu Cys Lys Ala Met Thr Leu

			180					185					190		
Cys	He	Cys	Tyr	Ala	Ala	Ser	Ile	Gly	Gly	Thr	Ala	Thr	Leu	Thr	Gly
		195					200					205			
Thr	Gly	Pro	Asn	Val	Val	Leu	Leu	Gly	Gln	Met	Asn	Glu	Leu	Phe	Pro
	210					215					220				
Asp	Ser	Lys	Asp	Leu	Val	Asn	Phe	Ala	Ser	Trp	Phe	Ala	Phe	Ala	Phe
225					230					235					240
Pro	Asn	Met	Leu	Val	Met	Leu	Leu	Phe	Ala	Trp	Leu	Trp	Leu	Gln	Phe
				245					250					255	
Val	Tyr	Met	Arg	Phe	Asn	Phe	Lys	Lys	Ser	Trp	Gly	Cys	Gly	Leu	Glu
			260					265					270		
Ser	Lys	Lys	Asn	Glu	Lys	Ala	Ala	Leu	Lys	Val	Leu	Gln	Glu	Glu	Tyr
		275					280					285			
Arg	Lys	Leu	Gly	Pro	Leu	Ser	Phe	Ala	Glu	11e	Asn	Val	Leu	Пе	Cys
	290					295					300				
Phe	Phe	Leu	Leu	Val	lle	Leu	Trp	Phe	Ser	Arg	Asp	Pro	Gly	Phe	Met
305					310					315					320
Pro	Gly	Trp	Leu	Thr	Val	Ala	Trp	Val	Glu	Gly	Glu	Thr	Lys		
				325					330						

<211> 451

<212> PRT

<213> Homo sapiens

<400> 4814

Met Asn Lys Ala Pro Gln Ser Thr Gly Pro Pro Pro Ala Pro Ser Pro

1 5 10 15

Gly Leu Pro Gln Pro Ala Phe Pro Pro Gly Gln Thr Ala Pro Val Val

20 25 30

Phe Ser Thr Pro Gln Ala Thr Gln Met Asn Thr Pro Ser Gln Pro Arg 35 40 45

Gln His Phe Tyr Pro Ser Arg Ala Gln Pro Pro Ser Ser Ala Ala Ser 50 55 60

Arg Val Gln Ser Ala Ala Pro Ala Arg Pro Gly Pro Ala Ala Leu Pro

65					70					75					80
Pro	Val	Ser	Pro	Leu	Lys	Ala	Λla	Leu	Ser	Glu	Glu	Glu	Leu	Glu	Lys
				85					90					95	
Lys	Ser	Lys	Ala	He	He	Glu	Glu	Tyr	Leu	His	Leu	Asn	Asp	Met	Lys
			100					105					110		
Glu	Ala	Val	Gln	Cys	Va]	Gln	Glu	Leu	Ala	Ser	Pro	Ser	Leu	Leu	Phe
		115					120					125			
lle	Phe	Val	Arg	His	Gly	Val	Glu	Ser	Thr	Leu	Glu	Arg	Ser	Ala	11e
	130					135					140				
Ala	Arg	Glu	His	Met	Gly	Gln	Leu	Leu	His	Gln	Leu	Leu	Cys	Ala	Gly
145					150					155					160
His	Leu	Ser	Thr	Ala	Gln	Tyr	Tyr	Gln	Gly	Leu	Tyr	Glu	lle	Leu	Glu
				165					170					175	
Leu	Ala	Glu	Λsp	Met	Glu	11e	Asp	He	Pro	His	Val	Trp	Leu	Tyr	Leu
			180					185					190		
Ala	Glu	Leu	Val	Thr	Pro	lle	Leu	Gln	Glu	Gly	Gly	Val	Pro	Met	Gly
		195					200					205			
Glu	Leu	Phe	Arg	Glu	Ile	Thr	Lys	Pro	Leu	Arg	Pro	Leu	Gly	Lys	Ala
	210					215					220				
Ala	Ser	Leu	Leu	Leu	Glu	He	Leu	Gly	Leu	Leu	Cys	Lys	Ser	Met	Gly.
225					230					235					240
Pro	Lys	Lys	Val	Gly	Thr	Leu	Trp	Arg	Glu	Ala	Gly	Leu	Ser	Trp	Lys
				245					250					255	
Glu	Phe	Leu	Pro	Glu	Gly	Gln	Asp	He	G1 y	Ala	Phe	Val	Ala	Glu	GIn
			260					265					270		
Lys	Val	Glu	Tyr	Thr	Leu	Gly	Glu	Glu	Ser	Glu	Ala	Pro	Gly	Gln	Arg
		275					280					285			
Ala	Leu	Pro	Ser	Glu	Glu	Leu	Asn	Arg	Gln	Leu	Glu	Lys	Leu	Leu	Lys
	290					295					300				
Glu	Gly	Ser	Ser	Asn	Gln	Arg	Val	Phe	Asp	Trp	He	Glu	Ala	Asn	Leu
305					310					315					320
Ser	Glu	GIn	Gln	He	Val	Ser	Asn	Thr	Leu	Val	Arg	Ala	Leu		Thr
				325					330					335	
Ala	Val	Cys			Ala	He	He		Glu	Thr	Pro	Leu			Asp
			340					345					350		
Val	Ala	Val	Leu	Lys	Ala	Arg	Ala	Lys	Leu	Leu	Gln	Lys	Tyr	Leu	Cys

Asp Glu Gln Lys Glu Leu Gln Ala Leu Tyr Ala Leu Gln Ala Leu Val Val Thr Leu Glu Gln Pro Pro Asn Leu Leu Arg Met Phe Phe Asp Ala Leu Tyr Asp Glu Asp Val Val Lys Glu Asp Ala Phe Tyr Ser Trp Glu Ser Ser Lys Asp Pro Ala Glu Gln Gln Gly Lys Gly Val Ala Leu Lys Ser Val Thr Ala Phe Phe Lys Trp Leu Arg Glu Ala Glu Glu Glu Ser Asp His Asn <210> 4815 <211> 694 <212> PRT <213> Homo sapiens <400> 4815 Met Glu Arg Ala Met Glu Gln Leu Asn Arg Leu Thr Arg Ser Leu Arg Arg Ala Arg Thr Val Glu Leu Pro Glu Asp Asn Glu Thr Ala Val Tyr Thr Leu Met Pro Met Val Met Ala Asp Gln His Arg Ser Val Ser Glu Leu Leu Ser Asn Ser Lys Phe Asp Val Asn Tyr Ala Phe Gly Arg Val Lvs Arg Ser Leu Leu His Ile Ala Ala Asn Cys Gly Ser Val Glu Cys Leu Val Leu Leu Lys Lys Gly Ala Asn Pro Asn Tyr Gln Asp Ile Ser Gly Cys Thr Pro Leu His Leu Ala Ala Arg Asn Gly Gln Lys Lys

Cys Met Ser Lys Leu Leu Glu Tyr Ser Ala Asp Val Asn lle Cys Asn

		115					120					125			
Asn	Glu	G1 y	Pro	Thr	Ala	He	His	Trp	Leu	Ala	Val	Asn	Gly	Arg	Thr
	130					135					140				
Glu	Leu	Leu	His	Asp	Leu	Val	G1n	His	Val	Ser	Asp	Va]	Asp	Val	Glu
145					150					155					160
Asp	Ala	Met	Gly	Gln	Thr	Ala	Leu	His	Val	Ala	Cys	Gln	Asn	Gly	His
				165					170					175	
Lys	Thr	Thr	Val	Gln	Cys	Leu	Leu	Asp	Ser	Gly	Ala	Asp	lle	Asn	Arg
			180					185					190		
Pro	Asn	Val	Ser	Gly	Ala	Thr	Pro	Leu	Tyr	Phe	Ala	Cys	Ser	His	Gly
		195					200					205			
Gln	Arg	Asp	Thr	Ala	Gln	lle	Leu	Leu	Leu	Arg	Gly	Ala	Lys	Tyr	Leu
	210					215					220				
Pro	Asp	Lys	Asn	Gly	Val	Thr	Pro	Leu	Asp	Leu	Cys	Val	Gln	Gly	Gly
225					230					235					240
Tyr	Gly	Glu	Thr	Cys	Glu	Val	Leu	Ile	Gln	Tyr	His	Pro	Arg	Leu	Phe
				245					250					255	
Gln	Thr	Ile	Ile	Gln	Met	Thr	Gln	Asn	Glu	Asp	Leu	Arg	Glu	Asn	Met
			260					265					270		
Leu	Arg	Gln	Val	Leu	Glu	His	Leu	Ser	Gln	Gln	Ser	Glu	Ser	Gln	Tyr
		275					280					285			
Leu	Lys	Ile	Leu	Thr	Ser	Leu	Ala	Glu	Val	Ser	Thr	Thr	Asn	Gly	His
	290					295					300				
Lys	Leu	Leu	Ser	Leu	Ser	Ser	Asn	Tyr	Asp	Ala	Gln	Met	Lys	Ser	Leu
305					310					315					320
Leu	Arg	He	Val	Arg	Met	Phe	Cys	His	Val	Phe	Arg	lle	Gly	Pro	Ser
				325					330					335	
Ser	Pro	Ser	Asn	Gly	lle	Asp	Met	Gly	Tyr	Asn	Gly	Asn	Lys	Thr	Pro
			340					345					350		
Arg	Ser	Gln	Val	Phe	Lys	Pro	Leu	G) u	Leu	Leu	Trp	His	Ser	Leu	Asp
		355					360					365			
Glu	Trp	Leu	Val	Leu	He	Ala	Thr	Glu	Leu	Met	Lys	Asn	Lys	Arg	Asp
	370					375					380				
Ser	Thr	Glu	He	Thr	Ser	He	Leu	Leu	Lys	Gln	Lys	Gly	Gln	Asp	Gln
385					390					395					400
Acn	Ala	Ala	Sor	116	Pro	Pro	Phe	Glu	Pro	Pro	G1v	Pro	G1v	Ser	Tyr

				405					410					415	
Glu	Λsn	Leu	Ser	Thr	Gly	Thr	Arg	Glu	Ser	Lys	Pro	Asp	Ala	Leu	Ala
			420					425					430		
Gly	Arg	Gln	Glu	Ala	Ser	Ala	Лѕр	Cys	Gln	Asp	Val	He	Ser	Met	Thr
		435					440					445			
Ala	Asn	Arg	Leu	Ser	Ala	Val	11e	Gln	Ala	Phe	Tyr	Met	Cys	Cys	Ser
	450					455					460				
Cys	Gln	Met	Pro	Pro	Gly	Met	Thr	Ser	Pro	Arg	Phe	Ile	Glu	Phe	Val
465					470					475					480
Cys	Lys	His	Asp	Glu	Val	Leu	Lys	Cys	Phe	Val	Asn	Arg	Asn	Pro	Lys
				485					490					495	
lle	lle	Phe	Asp	His	Phe	His	Phe	Leu	Leu	Glu	Cys	Pro	Glu	Leu	Met
			500					505					510		
Ser	Arg	Phe	Met	His	lle	He	Lys	Ala	Gln	Ala	Glu	Tyr	Val	Gln	Leu
		515					520					525			
Val	Thr	Glu	Leu	Ārg	Met	Thr	Arg	Ala	Ile	Gln	Pro	Gln	Tle	Asn	Ala
	530					535					540				
Phe	Leu	Gln	Gly	Phe	His	Met	Phe	Ile	Pro	Pro	Ser	Leu	Ile	Gln	Leu
545					550					555					560
Phe	Asp	Glu	Tyr	Glu	Leu	Glu	Leu	Leu	Leu	Ser	Gly	Met	Pro	Glu	He
				565					570					575	
Asp	Val	Ser	Asp	Trp	lle	Lys	Asn	Thr	Glu	Tyr	Thr	Ser	Gly	Tyr	Glu
			580					585					590		
Arg	Glu	Asp	Pro	Val	He	Gln	Trp	Phe	Trp	Glu	Val	Val	Glu	Asp	He
		595					600					605			
Thr	Gln	Glu	Glu	Arg	Val	Leu	Leu	Leu	Gln	Phe	Val	Thr	Gly	Ser	Ser
	610					615					620				
Arg	Val	Pro	His	Gly	Gly	Phe	Ala	Asn	He	Met	Gly	Gly	Ser	Gly	Leu
625					630					635					640
Gln	Asn	Phe	Thr	lle	Ala	Ala	Val	Pro	Tyr	Thr	Pro	Asn	Leu	Leu	Pro
				645					650					655	
Thr	Ser	Ser	Thr	Cys	He	Asn	Met	Leu	Lys	Leu	Pro	Glu	Tyr	Pro	Ser
			660					665					670		
Lys	Glu	11e	Leu	Lys	Asp	Arg	Leu	Leu	Val	Ala	Leu	His	Cys	Gly	Ser
		675					680					685			
Tyr	Gly	Tyr	Thr	Met	Ala										

690 .

<210> 4816

<211	> 64	19													
<212	?> PF	RT													
<213	8> Hc	omo s	sapie	ens											
<400)> 48	316													
Met	Pro	Lys	Ala	Glu	Ala	Asn	Leu	Gly	Gly	Leu	Ser	Ser	Leu	Val	Val
1				5					10					15	
Asn	Thr	Pro	He	Thr	Ser	Val	Ser	Leu	Ser	His	Ser	Ser	Ser	Glu	Ser
			20					25					30		
Ser	Lys	Met	Ser	Glu	Ser	Lys	Asp	Gln	Glu	Asn	Asn	Cys	Glu	Arg	Pro
		35					40					45			
Lys	Glu	Ser	Asn	Val	Leu	His	Pro	Asn	G1 y	Glu	Cys	Pro	Val	Lys	Ser
	50					55					60				
Glu	Pro	Thr	Glu	Pro	Gly	Asp	Glu	Asp	Glu	Glu	Asp	Ala	Tyr	Ser	Asn
65					70					75					80
Glu	Leu	Asp	Asp	Glu	Glu	Val	Leu	Gly	Glu	Leu	Thr	Asp	Ser	He	Gly
				85					90					95	
Asn	Lys	Asp	Phe	Pro	Leu	Leu	Asn	Gln	Ser	He	Ser	Pro	Leu	Ser	Ser
			100					105					110		
Ser	Val	Leu	Lys	Phe	He	Glu	Lys	Gly	Thr	Ser	Ser	Ser	Ser	Ala	Thr
		115					120					125			
Val	Ser	Asp	Asp	Thr	Glu	Lys	Lys	Lys	Gln	Thr	Ala	Ala	Val	Arg	Ala
	130					135					140				
Ser	Gly	Ser	Val	Ala	Ser	Asn	Tyr	Gly	He	Ser	Gly	Lys	Asp	Phe	Ala
145					150					155					160
Asp	Ala	Ser	Ala	Ser	Lys	Asp	Ser	Ala	Thr	Ala	Ala	His	Pro	Ser	Glu
				165					170					175	
lle	Ala	Arg	Gly	Asp	Glu	Asp	Ser	Ser	Ala	Thr	Pro	His	Gln	His	Gly
			180					185					190		
Phe	Thr	Pro	Ser	Thr	Pro	G1 y	Thr	Pro	Gly	Pro	G] y	Gly	Asp	Gly	Ser
		195					200					205			
Pro	G1y	Ser	Gly	lle	Glu	Cys	Pro	Lys	Cys	Asp	Thr	Val	Leu	Gly	Ser

	210					215					220				
Ser	Arg	Ser	Leu	G1 y	Gly	His	Met	Thr	Met	Met	His	Ser	Arg	Asn	Ser
225					230					235					240
Cys	Lys	Thr	Leu	Lys	Cys	Pro	Lys	Cys	Asn	Trp	His	Tyr	Lys	Tyr	Gln
				245					250					255	
Gln	Thr	Leu	Glu	Ala	His	Met	Lys	Glu	Lys	His	Pro	Glu	Pro	Gly	G1 y
			260					265					270		
Ser	Cys	Val	Tyr	Cys	Lys	Thr	Gly	Gln	Pro	His	Pro	Arg	Leu	Ala	Arg
		275					280					285			
Gly	Glu	Ser	Tyr	Thr	Cys	Gly	Tyr	Lys	Pro	Phe	Arg	Cys	Glu	Val	Cys
	290					295					300				
Asn	Tyr	Ser	Thr	Thr	Thr	Lys	Gly	Asn	Leu	Ser	Ile	His	Met	Gln	Ser
305					310					315					320
Asp	Lys	His	Leu	Asn	Asn	Val	Gln	Asn	Leu	Gln	Asn	Gly	Asn	Gly	Glu
				325					330					335	
Gln	Val	Phe	Gly	His	Ser	Ala	Pro	Ala	Pro	Asn	Thr	Ser	Leu	Ser	Gly
			340					345					350		
Cys	Gly	Thr	Pro	Ser	Pro	Ser	Lys	Pro	Lys	Gln	Lys		Thr	Trp	Arg
		355					360					365			
Cys		Val	Cys	Asp	Tyr		Thr	Asn	Val	Ala	Arg	Asn	Leu	Arg	He
	370					375					380				
	Met	Thr	Ser	Glu		His	Met	His	Asn		Met	Leu	Leu	GIn	
385					390					395					400
Asn	Met	Lys	GIn		GIn	His	Asn	Leu		Leu	Gly	Leu	Ala		Ala
0.1		61		405	0.1	T.	т	,	410	61		т 1	C1	415	т)
Glu	Ala	Glu			GIn	lyr	lyr			61n	Asn	11e		Leu	Inr
61	14 .	,	420		Λ	D	۸1.	425		C1	1	Mat	430	1 an	Duo
61 y	Met			GIU	Asn	Pro		Asp	Pro	GIN	Leu	мет 445	116	ASII	Pro
DL -	C1	435		Duo	Λlα	Tha	440	110	110	Lou	A10		Clv	Lou	Gly
rne		ren	ASP	FJO	на	455	ма	ита	ма	Leu	460	110	GIŸ	Leu	Oly
C I u	450	Sor	Dro	Tyr	110		Acn	Pro	Δ1a	Lou	Lys	Leu	Phe	Gln	Cvs
465		261	110	1 1/1	470		цар	110	та	475		LCu	1 110	0111	480
		Cve	Asn	lve			Ser	Asn	Ser		Glu	Ala	Leu	Ser	
ma	, CI J	Cys	7,511	485			501		490		014			495	
Hic	Val	Sor	Sor			Ser	Lou	Pro			Glu	Trn	Aro		Val

Ile Gly Asp Ile Tyr Gln Cys Lys Leu Cys Asn Tyr Asn Thr Gln Leu Lys Ala Asn Phe Gln Leu His Cys Glu Thr Asp Lys His Met Gln Lys Tyr Gln Leu Val Ala His Ile Lys Glu Gly Gly Lys Ser Asn Glu Trp Arg Leu Lys Cys Ile Ala Ile Gly Asn Pro Val His Leu Lys Cys Asn Ala Cys Asp Tyr Tyr Thr Asn Ser Val Asp Lys Leu Arg Leu His Thr Thr Asn His Arg His Glu Ala Ala Leu Lys Leu Tyr Lys Val Ser Ser Asp Ile His Phe Arg Trp His Arg Val Glu Lys Gly Ile Asn Ser Phe Arg Ala Trp Ser Thr Ser Leu Gln Leu Lys Glu Lys Lys Arg Glu Lys Thr Ser Lys Gly Arg Gly His Ser Phe <210> 4817 <211> 808 <212> PRT <213> Homo sapiens <400> 4817 Met Leu Glu Gly His Glu Ser Tyr Asp Thr Glu Asn Phe Tyr Phe Arg Glu Ile Arg Lys Asn Leu Gln Glu Val Asp Phe Gln Trp Lys Asp Gly Glu Ile Asn Tyr Lys Glu Gly Pro Met Thr His Lys Asn Asn Leu Thr

Gly Gln Arg Val Arg His Ser Gln Gly Asp Val Glu Asn Lys His Met

Glu	Asn	Gln	Leu	lle	Leu	Arg	Phe	Gln	Ser	Gly	Leu	Gly	Glu	Leu	Gln
65					70					75					80
Lys	Phe	Gln	Thr	Ala	Glu	Lys	11e	Tyr	G1y	Cys	Asn	Gln	He	Glu	Arg
				85					90					95	
Thr	Val	Asn	Asn	Cys	Phe	Leu	Ala	Ser	Pro	Leu	G1n	Arg	Пе	Phe	Pro
			100					105					110		
Gly	Val	Gln	Thr	Asn	lle	Ser	Arg	Lys	Tyr	Gly	Λsn	Asp	Phe	Leu	Gln
		115					120					125			
Leu	Ser	Leu	Pro	Thr	Gln	Asp	Glu	Lys	Thr	His	lle	Arg	Glu	Lys	Pro
	130					135					140				
Tyr	Ile	Gly	Asn	Glu	Cys	Gly	Lys	Ala	Phe	Arg	Val	Ser	Ser	Ser	Leu
145					150					155					160
He	Asn	His	Gln	Met	11e	llis	Thr	Thr	Glu	Lys	Pro	Tyr	Arg	Cys	Asn
				165					170					175	
Glu	Ser	Gly	Lys	Ala	Phe	His	Arg	Gly	Ser	Leu	Leu	Thr	Val	His	Gln
			180					185					190		
lle	Val	His	Thr	Arg	Gly	Lys	Pro	Tyr	Gln	Cys	Asp	Val	Cys	Gly	Arg
		195					200					205			
Ile	Phe	Arg	Gln	Asn	Ser	Asp	Leu	Val	Asn	His	Arg	Arg	Ser	His	Thr
	210					215					220				
Gly	Asp	Lys	Pro	Tyr	lle	Cys	Asn	Glu	Cys	G1 y	Lys	Ser	Phe	Ser	Lys
225					230					235					240
Ser	Ser	His	Leu	Ala	Val	His	G] n	Arg	lle	His	Thr	Gly	Glu	Lys	Pro
				245					250					255	
Tyr	Lys	Cys	Asn	Arg	Cys	Gly	Lys	Cys	Phe	Ser	Gln	Ser	Ser	Ser	Leu
			260					265					270		
Ala	Thr	His	Gln	Thr	Val	His	Thr	Gly	Asp	Lys	Pro	Tyr	Lys	Cys	Asn
		275					280					285			
Glu	Cys	Gly	Lys	Thr	Phe	Lys	Arg	Asn	Ser	Ser	Leu	Thr	Ala	His	His
	290					295					300				
He	He	His	Ala	Gly	Lys	Lys	Pro	Tyr	Thr	Cys	Asp	Val	Cys	Gly	Lys
305					310					315					320
Val	Phe	Tyr	Gln	Asn	Ser	Gln	Leu	Val	Arg	His	Gln	He	lle	His	Thr
				325					330					335	
Gly	Glu	Thr	Pro	Tyr	Lys	Cys	Asn	Glu	Cys	Gly	Lys	Val	Phe	Phe	Gln
			340					345					350		

Arg	Ser	Arg	Leu	Ala	Gly	His	Arg	Arg	Ile	His	Thr	G1 y	Glu	Lys	Pro
		355					360					365			
Tyr	Lys	Cys	Asn	Glu	Cys	Gly	Lys	Val	Phe	Ser	Gln	His	Ser	His	Leu
	370					375					380				
Ala	Val	His	Gln	Arg	Val	His	Thr	Gly	Glu	Lys	Pro	Tyr	Lys	Cys	Asn
385					390					395					400
Glu	Cys	Gly	Lys	Ala	Phe	Asn	Trp	Gly	Ser	Leu	Leu	Thr	Val	His	Gln
				405					410					415	
Arg	He	His	Thr	Gly	Glu	Lys	Pro	Tyr	Lys	Cys	Asn	Val	Cys	Gly	Lys
			420					425					430		
Val	Phe	Asn	Tyr	Gly	Gly	Tyr	Leu	Ser	Val	His	Met	Arg	Cys	His	Thr
		435					440					445			
Gly	Glu	Lys	Pro	Leu	His	Cys	Asn	Lys	Cys	Gly	Met	Val	Phe	Thr	Tyr
	450					455					460				
Tyr	Ser	Cys	Leu	Ala	Arg	His	Gln	Arg	Met	His	Thr	Gly	Glu	Lys	Pro
465					470					475					480
Tyr	Lys	Cys	Asn	Val	Cys	Gly	Lys	Val	Phe	lle	Asp	Ser	Gly	Asn	Leu
				485					490					495	
Ser	Ile	His	Arg	.Arg	Ser	His	Thr	Gly	Glu	Lys	Pro	Phe		Cys	Asn
			500					505					510		
Glu	Cys		Lys	Val	Phe	Ser		Tyr	Ser	Cys	Leu		Arg	His	Arg
		515					520	_				525	_	0.1	
Lys		His	Thr	Gly	Glu		Pro	Tyr	Lys	Cys		Asp	Cys	Gly	Lys
	530					535	_	<i>-</i>			540		7.3		(D)
	Tyr	Thr	G1n	Arg		Ser	Leu	Thr	Lys		Leu	Val	He	HIS	Thr
545	61		В	т	550	6	•	C1	DI	555	C1	41.	101	31.	560
Gly	61u	Asn	Pro			Cys	Asn	GIU			GIU	Ala	Pne		Gln
C	C	1	I	565		Т	ш	Λ	570		Thu	Clu	C1	575	Dwo
ser	ser	Lys		на	Arg	1 y r	nis	585	ASH	110	HIII	Giy	590	Lys	Pro
11.5 =	Lua	Cua	580	C1	Cua	C1	A 20 cz		Dha	Sor	llic	Lvc		Sor	Lou
HIS	Lys	595	Sei	Glu	Cys	Gly	600	1111	гне	261	1115	605		261	Leu
Va1	Tvr		Cln	Ana	Ara	Hic		Cl v	Clu	Mot	Pro			Cvs	He
val	610		0111	M g	пів	615	1111	01 y	oru	sie t	620	1 31	Lys	Oys	110
Glu			lve	Va1	Pho		Sor	Thr	Thr	Thr		Ala	Aro	Hic	Arg
625		OIY	Lys	· al	630		261	1111	1111	635		1110	11.1 g	1113	640
020					555					2.55					

Arg Ile His Thr Gly Glu Lys Pro Tyr Lys Cys Asn Glu Cys Gly Lys Val Phe Arg Tyr Arg Ser Gly Leu Ala Arg His Trp Ser Ile His Thr Gly Glu Lys Pro Tyr Lys Cys Asn Glu Cys Gly Lys Ala Phe Arg Val Arg Ser Ile Leu Leu Asn His Gln Met Met His Thr Gly Glu Lys Pro Tyr Lys Cys Asn Glu Cys Gly Lys Ala Phe Ile Glu Arg Ser Asn Leu Val Tyr His Gln Arg Asn His Thr Gly Glu Lys Pro Tyr Lys Cys Met Glu Cys Gly Lys Ala Phe Gly Arg Arg Ser Cys Leu Thr Lys His Gln Arg Tle His Ser Ser Glu Lys Pro Tyr Lys Cys Asn Glu Cys Gly Lys Ser Tyr Ile Ser Arg Ser Gly Leu Thr Lys His Gln Ile Lys His Ala Gly Glu Asn Leu Thr Thr Lys Leu Asn Val Glu Arg Pro Leu Asp Val Val Leu Thr Ser Gly Ile Pro Lys

<210> 4818

<211> 602

<212> PRT

<213> Homo sapiens

<400> 4818

Gly	Val 50	Cys	Leu	Leu	Tyr	Asp 55	Leu	Leu	His	Ser	Pro 60	Trp	Leu	Gln	Ala
	Leu	Lys	lle	Tyr	Asp	Cys	Leu	Gln	Glu		Lys	Glu	Lys	Lys	
65		4.7	T)	D	70	. 1	C1	12 1		75	т	C1	V 1	17 1	80
val	Pro	Ala	Ihr		His	Ala	GIN	Val		Ser	ıyr	Glu	val		61u
	,		C1	85 TI	D	TI	C	D	90	7.1	C.T.	C)		95	C1
Leu	Leu	Arg		Ihr	Pro	Inr	Ser		Glu	116	GIN	61 u		Arg	GIN
		0.1	100	D		DI		105		T)	7.1		110		C1
Met	Leu		Ala	Pro	His	Phe		GIY	Ala	Ihr	116		Arg	HIS	Glü
	mı.	115					120		~ 1			125	0.1		
Met		Gly	Asp	He	Leu		Ala	Arg	He	He		Gly	G1 y	Leu	Ala
	130	_		_		135					140				
	Arg	Ser	Gly	Leu	Leu	Tyr	Ala	Gly	Asp		Leu	Val	Glu	Val	
145					150					155					160
Gly	Val	Ser	Val		Gly	Leu	Asp	Pro		Gln	Val	He	His	He	Leu
				165					170					175	
Ala	Met	Ser		Gly	Thr	He	Met		Lys	Val	Val	Pro		Ser	Asp
			180					185					190		
Pro	Pro		Asn	Ser	Gln	Gln	Met	Val	Tyr	Val	Arg		Met	Thr	Glu
		195					200					205			
Tyr	Trp	Pro	Gln	Glu	Asp	Pro	Asp	lle	Pro	Cys	Met	Asp	Ala	Gl y	Leu
	210					215					220				
Pro	Phe	Gln	Lys	Gly	Asp	He	Leu	Gln	He	Val	Asp	Gln	Asn	Asp	Ala
225					230					235					240
Leu	Trp	Trp	G1n	Ala	Arg	Lys	He	Ser	Asp	Pro	Ala	Thr	Cys	Ala	Gly
				245					250					255	
Leu	Va]	Pro	Ser	Asn	His	Leu	Leu	Lys	Arg	Trp	Ser	Phe	Ala	Leu	Val
			260					265					270		
Ala	Gln	Ala	Gly	Val	Gln	Trp	His	Tvr	Leu	Asp	Ser	Leu _.	Gln	Pro	Leu
		275					280					285			
Pro	Pro	Gly	Phe	Lys	Arg	Phe	Ser	Cys	Leu	Ser	Leu	Pro	Arg	Ser	Trp
	290					295					300				
Asp	Tyr	He	Glu	Asp	Asp	Met	Lys	He	Asp	Glu	Lys	Cys	Val	Glu	Ala
305					310					315					320
Asp	Glu	Glu	Thr	Phe	Glu	Ser	Asp	Lys	Glu	Glu	Phe	Val	Gly	Tyr	Gly
				325					330					335	

Gln	Lys	Phe	Phe	He	Ala	Gly	Phe	Arg	Arg	Ser	Met	Arg	Leu	Cys	Arg
			340					345					350		
Arg	Lys	Ser	His	Leu	Ser	Pro	Leu	His	Ala	Ser	Val	Cys	Cys	Thr	Gly
		355					360					365			
Ser	Cys	Tyr	Ser	Ala	Val	Gly	Ala	Pro	Tyr	Glu	Glu	Val	Val	Arg	Tyr
	370					375					380				
Gln	Arg	Arg	Pro	Ser	Asp	Lys	Tyr	Arg	Leu	Ile	Val	Leu	Met	G1 y	Pro
385					390					395					400
Ser	Gly	Val	Gly	Val	Asn	Glu	Leu	Arg	Arg	Gln	Leu	Ile	Glu	Phe	Asn
				405					410					415	
Pro	Ser	His	Phe	Gln	Ser	Ala	Val	Pro	His	Thr	Thr	Arg	Thr	Lys	Lys
			420					425					430		
Ser	Tyr	Glu	Met	Asn	Gly	Arg	Glu	Tyr	His	Tyr	Val	Ser	Lys	Glu	Thr
		435					440					445			
Phe	Glu	Asn	Leu	He	Tyr	Ser	His	Arg	Met	Leu	Glu	Tyr	Gly	Glu	Tyr
	450					455					460				
Lys	Gly	His	Leu	Tyr	Gly	Thr	Ser	Val	Asp	Ala	Val	Gln	Thr	Val	Leu
465					470					475					480
Val	Glu	Gly	Lys	lle	Cys	Val	Met	Asp	Leu	Glu	Pro	Gln	Asp	Ile	Gln
				485					490					495	
Gly	Val	Arg	Thr	His	Glu	Leu	Lys	Pro	Tyr	Val	He	Phe	He	Lys	Pro
			500					505					510		
Ser	Asn	Met	Arg	Cys	Met	Lys	Gln	Ser	Arg	Lys	Asn	Ala	Lys	Val	11e
		515					520					525			
Thr	Asp	Tyr	Tyr	Val	Asp	Met	Lys	Phe	Lys	Asp	Glu	Asp	Leu	Gln	Glu
	530					535					540				
Met	Glu	Asn	Leu	Ala	Gln	Arg	Met	Glu	Thr	Gln	Phe	Gly	Gln	Phe	Phe
545					550					555					560
Asp	His	Val	He	Val	Asn	Asp	Ser	Leu	His	Asp	Ala	Cys	Ala	Gln	Leu
				565					570					575	
Leu	Ser	Ala	Пе	G1n	Lys	Ala	Gln	Glu	Glu	Pro	Gln	Trp	Val	Pro	Ala
			580					585					590		
Thr	Trp	He	Ser	Ser	Asp	Thr	Glu	Ser	Gln						
		595					600								

```
<210> 4819
<211> 498
<212> PRT
<213> Homo sapiens
<400> 4819
Met Glu Ser Pro Arg Gly Trp Thr Leu Gln Val Ala Pro Glu Glu Gly
Gln Val Leu Cys Asn Val Lys Thr Ala Thr Arg Gly Leu Ser Glu Gly
                                                      30
             20
                                 25
Ala Val Ser Gly Gly Trp Gly Ala Trp Glu Asn Ser Thr Glu Val Pro
                             40
Arg Glu Ala Gly Asp Gly Gln Arg Gln Gln Ala Thr Leu Gly Ala Ala
                         55
Asp Glu Gln Gly Gly Pro Gly Arg Glu Leu Gly Pro Ala Asp Gly Gly
                     70
                                          75
65
Arg Asp Gly Ala Gly Pro Arg Ser Glu Pro Ala Asp Arg Ala Leu Arg
                                     90
                 85
Pro Ser Pro Leu Pro Glu Glu Pro Gly Cys Arg Cys Gly Glu Cys Gly
                                105
            100
Lys Ala Phe Ser Gln Gly Ser Tyr Leu Leu Gln His Arg Arg Val His
                            120
                                                 125
Thr Gly Glu Lys Pro Tyr Thr Cys Pro Glu Cys Gly Lys Ala Phe Ala
    130
                        135
Trp Ser Ser Asn Leu Ser Gln His Gln Arg lle His Ser Gly Glu Lys
                    150
                                         155
Pro Tyr Ala Cys Arg Glu Cys Gly Lys Ala Phe Arg Ala Gln Ser Gln
                                     170
                                                         175
                165
Leu 11c His His Gln Glu Thr His Ser Gly Leu Lys Pro Phe Arg Cys
            180
                                 185
Pro Asp Cys Gly Lys Ser Phe Gly Arg Ser Thr Thr Leu Val Gln His
                            200
                                                 205
Arg Arg Thr His Thr Gly Glu Lys Pro Tyr Glu Cys Pro Glu Cys Gly
                        215
                                             220
```

Lys Ala Phe Ser Trp Asn Ser Asn Phe Leu Glu His Arg Arg Val His

225					230					235					240
Thr	Gly	Ala	Arg	Pro	His	Ala	Cys	Arg	Asp	Cys	Gly	Lys	Ala	Phe	Ser
				245					250					255	
Gln	Ser	Ser	Asn	Leu	Ala	Glu	His	Leu	Lys	He	His	Ala	Gly	Ala	Arg
			260					265					270		
Pro	His	Ala	Cys	Pro	Asp	Cys	Gly	Lys	Ala	Phe	Val	Arg	Val	Ala	Gly
		275					280					285			
Leu	Arg	Gln	His	Arg	Arg	Thr	His	Ser	Ser	Glu	Lys	Pro	Phe	Pro	Cys
	290					295					300				
Ala	Glu	Cys	Gly	Lys	Ala	Phe	Λrg	Glu	Ser	Ser	Gln	Leu	Leu	Gln	His
305					310					315					320
Gln	Arg	Thr	His	Thr	Gly	Glu	Arg	Pro	Phe	G] u	Cys	Ala	Glu	Cys	Gly
				325					330					335	
Gln	Ala	Phe	Val	Met	Gly	Ser	Tyr	Leu	Ala	Glu	His	Arg	Arg	Val	His
			340					345					350		
Thr	Gly	Glu	Lys	Pro	His	Ala	Cys	Ala	Gln	Cys	Gly	Lys	Ala	Phe	Ser
		355					360					365			
Gln	Arg	Ser	Asn	Leu	Leu	Ser	His	Arg	Arg	Thr	His	Ser	Gly	Ala	Lys
	370					375					380				
Pro	Phe	Ala	Cys	Ala	Asp	Cys	Gly	Lys	Ala	Phe	Arg	Gly	Ser	Ser	G1 y
385					390					395					400
Leu	Ala	His	His	Arg	Leu	Ser	His	Thr	Gly	Glu	Arg	Pro	Phe	Ala	Cys
				405					410					415	
Ala	Glu	Cys	Gl y	Lys	Ala	Phe	Arg	Gly	Ser	Ser	Glu	Leu	Λrg	Gln	His
			420					425					430		
Gln	Arg	Leu	His	Ser	Gly	Glu	Arg	Pro	Phe	Val	Cys	Ala	His	Cys	Ser
		435					440					445			
Lys	Ala	Phe	Val	Arg	Lys	Ser	Glu	Leu	Leu	Ser	His	Arg	Arg	Thr	His
	450					455					460				
Thr	Gly	Glu	۸rg	Pro	Tyr	Ala	Cys	Gly	Glu	Cys	Gly	Lys	Pro	Phe	Ser
465					470					475					480
His	Arg	Cys	Asn	Leu	Asn	Glu	His	Gln	Lys	Arg	His	Gly	Gly	Arg	Ala
				485					490					495	
Ala	Pro														

```
<210> 4820
<211> 486
<212> PRT
<213> Homo sapiens
```

<400> 443 Met Arg Gly Arg Gly Ser Gln Gln Gln Gln Pro Thr Arg Arg Gln Gly Gln Lys Leu Pro Ser Pro Ser Pro Ala Gly Lys Tyr Glu Ser Ala Gln Pro Gly Gly Thr Gln Pro Glu Pro Gly Leu Gly Ala Arg Met Ala Ile His Lys Ala Leu Val Met Cys Leu Gly Leu Pro Leu Phe Leu Phe Pro Gly Ala Trp Ala Gln Gly His Val Pro Pro Gly Cys Ser Gln Gly Leu Asn Pro Leu Tyr Tyr Asn Leu Cys Asp Arg Ser Gly Ala Trp Gly Ile Val Leu Glu Ala Val Ala Gly Ala Gly Ile Val Thr Thr Phe Val Leu Thr Ile Ile Leu Val Ala Ser Leu Pro Phe Val Gln Asp Thr Lys Lys Arg Ser Leu Leu Gly Thr Gln Val Phe Phe Leu Leu Gly Thr Leu Gly Leu Phe Cys Leu Val Phe Ala Cys Val Val Lys Pro Asp Phe Ser Thr Cys Ala Ser Arg Arg Phe Leu Phe Gly Val Leu Phe Ala Ile Cys Phe Ser Cys Leu Ala Ala Ilis Val Phe Ala Leu Asn Phe Leu Ala Arg Lys Asn His Gly Pro Arg Gly Trp Val 11e Phe Thr Val Ala Leu Leu Leu Thr Leu Val Glu Val Ile Ile Asn Thr Glu Trp Leu Ile Ile Thr Leu

Val	Arg	Gly	Ser	Gly	Glu	Gly	Gly	Pro	Gln	Gly	Asn	Ser	Ser	Ala	Gly
225					230					235					240
Trp	Ala	Val	Λla	Ser	Pro	Cys	Ala	Ile	Ala	۸sn	Met	Asp	Phe	Val	Met
				245					250					255	
Ala	Leu	He	Tyr	Val	Met	Leu	Leu	Leu	Leu	Gly	Ala	Phe	Leu	Gly	Ala
			260					265					270		
Trp	Pro	Ala	Leu	Cys	Gly	Arg	Tyr	Lys	Arg	Trp	Arg	Lys	His	Gly	Val
		275		٠			280					285			
Phe	Val	Leu	Leu	Thr	Thr	Ala	Thr	Ser	Val	Ala	Ile	Trp	Val	Val	Trp
	290					295					300				
Ile	Val	Met	Tyr	Thr	Tyr	Gly	Asn	Lys	Gln	His	Asn	Ser	Pro	Thr	Trp
305					310					315					320
Asp	Asp	Pro	Thr	Leu	Ala	11e	Ala	Leu	Ala	Ala	Asn	Ala	Trp	Ala	Phe
				325					330					335	
Val	Leu	Phe	Tyr	Va]	He	Pro	Glu	Val	Ser	Gln	Va]	Thr	Lys	Ser	Ser
			340					345					350		
Pro	Glu	Gln	Ser	Tyr	Gln	Gly	Asp	Met	Tyr	Pro	Thr	Arg	Gly	.Val	Gly
		355					360					365			
Tyr	Glu	Thr	Ile	Leu	Lys	Glu	Gln	Lys	Gly	Gln	Ser	Met	Phe	Val	Glu
	370					375					380				
Asn	Lys	Ala	Phe	Ser	Met	Asp	Glu	Pro	Val	Ala	Ala	Lys	Arg	Pro	Val
385					390					395					400
Ser	Pro	Tyr	Ser	Gly	Tyr	Asn	Gly	Gln	Leu	Leu	Thr	Ser	Val	Tyr	Gln
				405					410					415	
Pro	Thr	Glu	Met	Ala	Leu	Met	His	Lys	Val	Pro	Ser	Glu	Gly	Ala	Tyr
			420					425					430		
Asp	lle	He	Leu	Pro	Arg	Ala	Thr	Ala	Asn	Ser	Gln		Met	Gły	Ser
		435					440					445			
Ala		Ser	Thr	Leu	Arg		Glu	Asp	Met	Tyr		Ala	Gln	Ser	His
	450					455					460				
	Ala	Ala	Thr	Pro			Asp	Gly	Lys		Ser	Gln	Va]	Phe	
465					470					475			•		480
Asn	Pro	Tyr	Val	Trp	Asp										

```
<211> 301
<212> PRT
<213> Homo sapiens
<400> 444
Met Tyr Thr Arg Arg Tyr Ser Ser Ile Ser Ser Thr Ile Met Asp Val
                                     10
Asp Ser Thr Ile Ser Ser Gly Arg Ser Thr Pro Ala Met Met Asn Gly
                                                      30
                                 25
Gln Gly Ser Thr Thr Ser Ser Ser Lys Asn Ile Ala Tyr Asn Cys Cys
                             40
Trp Asp Gln Cys Gln Ala Cys Phe Asn Ser Ser Pro Asp Leu Ala Asp
     50
                         55
                                              60
His Ile Arg Ser 11e His Val Asp Gly Gln Arg Gly Gly Val Phe Val
                     70
                                          75
Cys Leu Trp Lys Gly Cys Lys Val Tyr Asn Thr Pro Ser Thr Ser Gln
                                      90
                 85
Ser Trp Leu Gln Arg His Met Leu Thr His Ser Gly Asp Lys Pro Phe
            100
Lys Cys Val Val Gly Gly Cys Asn Ala Ser Phe Ala Ser Gln Gly Gly
                            120
Leu Ala Arg His Val Pro Thr His Phe Ser Gln Gln Asn Ser Ser Lys
                        135
                                             140
Val Ser Ser Gln Pro Lys Ala Lys Glu Glu Ser Pro Ser Lys Ala Gly
                    150
                                         155
                                                             160
145
Met Asn Lys Arg Arg Lys Leu Lys Asn Lys Arg Arg Arg Ser Leu Pro
                165
                                     170
Arg Pro His Asp Phe Phe Asp Ala Gln Thr Leu Asp Ala Ile Arg His
                                                     190
            180
                                 185
Arg Ala Ile Cys Phe Asn Leu Ser Ala His Ile Glu Ser Leu Gly Lys
                            200
Gly His Ser Val Val Phe His Ser Thr Val Ile Ala Lys Arg Lys Glu
                        215
                                             220
    210
Asp Ser Gly Lys Ile Lys Leu Leu Leu His Trp Met Pro Glu Asp Ile
```

 Leu Pro Asp Val
 Trp Val Asn Glu Ser Glu Arg His Gln Leu Lys Thr

 245
 250

 Lys Val Val His Leu Ser Lys Leu Pro Lys Asp Thr Ala Leu Leu 260
 265

 Asp Pro Asn 1le Tyr Arg Thr Met Pro Gln Lys Arg Leu Lys Arg Thr
 280

 Leu 1le Arg Lys Val Phe Asn Leu Tyr Leu Ser Lys Gln
 300

<210> 4822

<211> 626

<212> PRT

<213> Homo sapiens

<400> 445

Met Ala Val Ser Gly Phe Thr Leu Gly Thr Cys Ile Leu Leu Leu His 1 5 10 15 Ile Ser Tyr Val Ala Asn Tyr Pro Asn Gly Lys Val Thr Gln Ser Cys 20 25 30

His Gly Met Ile Pro Glu His Gly His Ser Pro Gln Ser Val Pro Val
35 40 45

His Asp lle Tyr Val Ser Gln Met Thr Phe Arg Pro Gly Asp Gln lle 50 55 60

Glu Val Thr Leu Ser Gly His Pro Phe Lys Gly Phe Leu Leu Glu Ala 65 70 75 80

Arg Asn Ala Glu Asp Leu Asn Gly Pro Pro Ile Gly Ser Phe Thr Leu 85 90 95

Ile Asp Ser Glu Val Ser Gln Leu Leu Thr Cys Glu Asp 11e Gln Gly
100 105 110

Ser Ala Val Ser His Arg Ser Ala Ser Lys Lys Thr Glu Ile Lys Val

Tyr Trp Asn Ala Pro Ser Ser Ala Pro Asn His Thr Gln Phe Leu Val 130 135 140

Thr Val Val Glu Lys Tyr Lys lle Tyr Trp Val Lys lle Pro Gly Pro 145 150 155 160

He	Ile	Ser	Gln	Pro	Asn	Ala	Phe	Pro	Phe	Thr	Thr	Pro	Lys	Ala	Thr
				165					170					175	
Val	Val	Leu	Leu	Pro	Thr	Leu	Pro	Pro	Val	Ser	His	Leu	Thr	Lys	Pro
			180					185					190		
Phe	Ser	Ala	Ser	Asp	Cys	Gly	Asn	Lys	Lys	Phe	Cys	Ile	Arg	Ser	Pro
		195					200					205			
Leu	Asn	Cys	Asp	Pro	Glu	Lys	Glu	Ala	Ser	Cys	Val	Phe	Leu	Ser	Phe
	210					215					220				
Thr	Arg	Asp	Asp	Gln	Ser	Val	Met	Val	Glu	Met	Ser	Gly	Pro	Ser	Lys
225					230					235					240
Gly	Tyr	Leu	Ser	Phe	Ala	Leu	Ser	His	Asp	Gln	Trp	Met	Gly	Asp	Asp
				245					250					255	
Asp	Ala	Tyr	Leu	Cys	11e	His	Glu	Asp	Gln	Thr	Val	Tyr	He	Gln	Pro
			260					265					270		
Ser	His	Leu	Thr	G1y	Arg	Ser	His	Pro	Val	Met	Asp	Ser	Arg	Asp	Thr
		275					280					285			
Leu	Glu	Asp	Met	Ala	Trp	Arg	Leu	Ala	Asp	Gly	Val	Met	Gln	Cys	Ser
	290					295					300				
Phe	Arg	Arg	Asn	lle	Thr	Leu	Pro	Gly	Val	Lys	Asn	Arg	Phe	Asp	Leu
305					310					315					320
Asn	Thr	Ser	Tyr	Tyr	He	Phe	Leu	Ala	Asp	Gly	Ala	Ala	Asn	Asp	Gly
				325					330					335	
Arg	Ile	Tyr	Lys	His	Ser	Gln	Gln	Pro	Leu	He	Thr	Tyr	Glu	Lys	Tyr
			340					345					350		
Asp	Val	Thr	Asp	Ser	Pro	Lys	Asn	He	Gly	Gly	Ser	His	Ser	Val	Leu
		355					360					365			
Leu	Leu	Lys	Val	His	Gly	Ala	Leu	Met	Phe	Val	Ala	Trp	Met	Thr	Thr
	370					375					380				
Val	Ser	He	Gly	Val	Leu	Val	Ala	Arg	Phe	Phe	Lys	Pro	Val	Trp	Ser
385					390					395					400
Lys	Ala	Phe	Leu	Leu	Gly	Glu	Ala	Ala	Trp	Phe	Gln	Val	His	Arg	Met
				405					410					415	
Leu	Met	Phe	Thr	Thr	Thr	Val	Leu	Thr	Cys	He	Ala	Phe	Val	Met	Pro
			420					425					430		
Phe	Ile	Tyr	Arg	Gly	Gly	Trp	Ser	Arg	His	Ala	Gly	Tyr	His	Pro	Tyr
		435					440					445			

Leu Gly Cys Ile Val Met Thr Leu Ala Val Leu Gln Pro Leu Leu Ala Val Phe Arg Pro Pro Leu His Asp Pro Arg Arg Gln Met Phe Asn Trp Thr His Trp Ser Met Gly Thr Ala Ala Arg Ile Ile Ala Val Ala Ala Met Phe Leu Gly Met Asp Leu Pro Gly Leu Asn Leu Pro Asp Ser Trp Lys Thr Tyr Ala Met Thr Gly Phe Val Ala Trp His Val Gly Thr Glu Val Val Leu Glu Leu Lys Tyr Trp Met Met Thr Glu Phe Arg Ser Phe Ser His Leu Leu Gln Trp Lys Gln Arg Val Met Leu Leu Lys Arg Gln Cys Trp Gln Phe Met Ser Val Gly Met Leu Leu Phe Ser Ser Tyr Phe Tyr Leu Gln Ser Thr Ile Tyr Glu Gln Ala Lys Thr Leu Ala Phe Ala Gly Gln Val Ile Ile Ile Ile Lys Pro Lys Lys Leu Glu Ala Cys Pro Asp Cys Leu Glu His Ile Cys Glu Phe Ser Leu Gly Arg Leu Gly Ser Cys Leu <210> 4823

<400> 446

<213> Homo sapiens

<211> 827 <212> PRT

Met lle Arg Arg Leu Lys Thr Glu Val Leu Thr Gln Leu Pro Pro Lys

1 5 10 15

Val Arg Gln Arg Ile Pro Phe Asp Leu Pro Ser Ala Ala Ala Lys Glu
20 25 30

Leu	Asn	Thr	Ser	Phe	Glu	Glu	Trp	Glu	Lys	Ile	Met	Arg	Thr	Pro	Asn
		35					40					45			
Ser	Gly	Ala	Met	Glu	Thr	Val	Met	Gly	Leu	lle	Thr	Arg	Met	Phe	Lys
	50					55					60				
Gln	Thr	Ala	He	Λla	Lys	Ala	Gly	Ala	Val	Lys	Asp	Tyr	lle	Lys	Met
65					70					75					80
Met	Leu	G1n	۸sn	Asp	Ser	Leu	Lys	Phe	Leu	Val	Phe	Ala	His	His	Leu
				85					90					95	
Ser	Met	Leu	Gln	Ala	Cys	Thr	Glu	Ala	Val	Ile	Glu	Asn	Lys	Thr	Arg
			100					105					110		
Tyr	Ile	Arg	Ile	Asp	G1 y	Ser	Val	Ser	Ser	Ser	Glu	Arg	lle	His	Leu
		115					120					125			
Val	Asn	Gln	Phe	Gln	Lys	Asp	Pro	Asp	Thr	Arg	Va]	Ala	He	Leu	Ser
	130					135					140				
lle	Gln	Ala	Ala	Gly	Gln	Gly	Leu	Thr	Phe	Thr	Ala	Ala	Ser	His	Val
145					150					155					160
Val	Phe	Ala	Glu	Leu	Tyr	Trp	Asp	Pro	Gly	His	He	Lys	Gln	Ala	Glu
				165					170					175	
Asp	Arg	Ala	His	Arg	Ile	Gly	Gln	Cys	Ser	Ser	Val	Asn	Ile	His	Tyr
			180					185					190		
Leu	He	Ala	Asn	Gly	Thṛ	Leu	Asp	Thr	Leu	Met	Trp	Gly	Met	Leu	Asn
		195					200					205			
Arg	Lys	Ala	Gln	Val	Thr	Gly	Ser	Thr	Leu	Asn	Gly	Arg	Lys	Glu	Lys
	210					215					220				
11e	Gln	Ala	Glu	Glu	Gly	Asp	Lys	Glu	Lys	Trp	Asp	Phe	Leu	Gln	Phe
225					230					235					240
Ala	Glu	Ala	Trp	Thr	Pro	Asn	Asp	Ser	Ser	Glu	Glu	Leu	Arg	Lys	Glu
				245					250					255	
Ala	Leu	Phe	Thr	His	Phe	Glu	Lys	Glu	Lys	Gln	His	Asp	lle	Arg	Ser
			260					265					270		
Phe	Phe	Val	Pro	Gln	Pro	Lys	Lys	Arg	Gln	Leu	Met	Thr	Ser	Cys	Asp
		275					280					285			
Glu	Ser	Lys	Arg	Phe	Arg	Glu	Glu	Asn	Thr	Val	Val	Ser	Ser	Asp	Pro
	290					295					300				
Thr	Lys	Thr	Ala	Ala	Arg	Asp	lle	lle	Asp	Tyr	Glu	Ser	Asp	Val	Glu
305					310					315					320

Pro	Glu	Thr	Lys	Arg	Leu	Lys	Leu	Ala	Ala	Ser	Glu	Asp	His	Cys	Ser
				325					330					335	
Pro	Ser	Ģlu	Glu	Thr	Pro	Ser	Gln	Ser	Lys	Gln	He	Arg	Thr	Pro	Leu
			340					345					350		
Val	Glu	Ser	Val	Gln	Glu	Ala	Lys	Ala	Gln	Leu	Thr	Thr	Pro	Ala	Phe
		355					360					365			
Pro	Val	Glu	Gly	Trp	Gln	Cys	Ser	Leu	Cys	Thr	Tyr	lle	Asn	Asn	Ser
	370					375					380				
Glu	Leu	Pro	Tyr	Cys	Glu	Met	Cys	Glu	Thr	Pro	Gln	Gly	Ser	Ala	Val
385					390					395					400
Met	Gln	lle	Asp	Ser	Leu	Asn	His	Ile	Gln	Asp	Lys	Asn	Glu	Lys	Asp
				405					410					415	
Asp	Ser	Gln	Lys	Asp	Thr	Ser	Lys	Lys	Val	G1n	Thr	lle	Ser	Asp	Cys
			420					425					430		
Glu	Lys	Gln	Ala	Leu	Ala	Gln	Ser	Glu	Pro	Gly	Gln	Leu	Ala	Asp	Ser
		435					440					445			
Lys	Glu	Glu	Thr	Pro	Lys	Ile	Glu	Lys	Glu	Asp	Gly	Leu	Thr	Ser	Gln
	450					455					460				
Pro	Gly	Asn	Glu	Gln	Trp	Lys	Ser	Ser	Asp	Thr	Leu	Pro	Val	Tyr	Asp
465					470					475					480
Thr	Leu	Met	Phe	Cys	Ala	Ser	Arg	Asn	Thr	Asp	Arg	lle	His	lle	Tyr
				485					490					495	
Thr	Lys	Asp	Gly	Lys	Gln	Met	Ser	Cys	Asn	Phe	He	Pro	Leu	Asp	lle
			500					505					510		
Lys	Leu	Asp	Leu	Trp	Glu	Asp	Leu	Pro	Ala	Ser	Phe	Gln	Leu	Lys	Gln
		515					520					525			
Tyr	Arg	Ser	Leu	lle	Leu	Arg	Phe	Val	Arg	Glu	Trp	Ser	Ser	Leu	Thr
	530					535					540				
	Met	Lys	Gln	Arg		He	Arg	Lys	Ser		Gln	Leu	Phe	Cys	
545					550					555					560
Pro	He	Leu	Ala		Glu	Glu	He	Thr		GIn	GIn	Thr	Lys	Gln	Asn
				565					570					575	
Cys	Thr	Lys		Tyr	He	Thr	Lys		Asp	Val	Ala	Val		Ser	Met
	,		580		., .	0.1	0.3	585	,		•	, ,	590		0.1
Asp	Lys	Ala	Lys	Asn	Val	Gly	Gly	His	Val	Arg	Leu	11e	lhr	Lys	61u

		595					600					605			
Ser	Arg	Pro	Arg	Asp	Pro	Phe	Thr	Lys	Lys	Leu	Leu	Glu	Asp	Gly	Ala
	610					615					620				
Cys	Val	Pro	Phe	Leu	Asn	Pro	Tyr	Thr	Val	Gln	Ala	Asp	Leu	Thr	Val
625					630					635					640
Lys	Pro	Ser	Thr	Ser	Lys	Gly	Tyr	Leu	Gln	Ala	Val	Asp	Asn	Glu	Gly
				645					650					655	
Asn	Pro	Leu	Cys	Leu	Arg	Cys	Gln	Gln	Pro	Thr	Cys	Gln	Thr	Lys	Gln
			660					665					670		
Ala	Cys	Lys	Ala	Asn	Ser	Trp	Asp	Ser	Arg	Phe	Cys	Ser	Leu	Lys	Cys
		675					680					685			
Gln	Glu	Glu	Phe	Trp	lle	Arg	Ser	Asn	Asn	Ser	Tyr	Leu	Arg	Ala	Lys
	690					695					700				
Val	Phe	Glu	Thr	Glu	His	Gly	Val	Cys	Gln	Leu	Cys	Asn	Val	Asn	Ala
705					710					715					720
Gln	Glu	Leu	Phe	Leu	Arg	Leu	Arg	Asp	Ala	Pro	Lys	Ser	Gln	Arg	Lys
				725					730					735	
Asn	Leu	Leu	Tyr	Ala	Thr	Trp	Thr	Ser	Lys	Leu	Pro	Leu	Glu	Gln	Leu
			740					745					750		
Asn	Glu	Met	lle	Arg	Asn	Pro	Gly	Glu	Gly	His	Phe	Trp	Gln	Val	Asp
		755					760					765			
His	He	Lys	Pro	Val	Tyr	Gly	Gly	Gly	Gly	Gln	Cys	Ser	Leu	Asp	Λsn
	770					775					780				
Leu	Gln	Thr	Leu	Cys	Thr	Val	Cys	His	Lys	Glu	Arg	Thr	Ala	Arg	Gln
785					790					795					800
Ala	Lys	Glu	Arg	Ser	Gln	Val	Arg	Arg	Gln	Ser	Leu	Ala	Ser	Lys	His
				805					810					815	
Gly	Ser	Asp	Ile	Thr	Arg	Phe	Leu	Val	Lys	Lys					
			820					825							

<211> 994

<212> PRT

<213≻ Homo sapiens

<400)> 44	17													
Met	Ser	Gln	Trp	Thr	Pro	Glu	Tyr	Lys	Glu	Leu	Tyr	Thr	Leu	Lys	Val
1				5					10					15	
Asp	Met	Lys	Ser	Glu	Ile	Pro	Ser	Asp	Ala	Pro	Lys	Thr	Gln	Glu	Ser
			20					25					30		
Leu	Lys	Gly	He	Leu	Leu	His	Pro	Glu	Pro	lle	Gly	Ala	Ala	Lys	Ser
		35					40					45			
Phe	Pro	Ala	Gly	Val	Glu	Met	Ile	Asn	Ser	Lys	Val	Gly	Asn	Glu	Phe
	50					55					60				
Ser	His	Leu	Cys	Asp	Asp	Ser	Gln	Lys	Gln	Glu	Lys	Glu	Met	Asn	G1 y
65					70					75					80
Asn	Gln	Gln	Glu	Gln	Glu	Lys	Ser	Leu	Val	Val	Arg	Lys	Lys	Arg	Lys
				85					90					95	
Ser	Gln	Gln	Ala	Gly	Pro	Ser	Tyr	Val	Gln	Asn	Cys	Val	Lys	Glu	Asn
			100					105					110		
Gln	Gly		Leu	Gly	Leu	Arg	Gln	His	Leu	Gly	Thr	Pro	Ser	Asp	Glu
		115					120					125			
Asp	Asn	Asp	Ser	Ser	Phe	Ser	Asp	Cys	Leu	Ser	Ser	Pro	Ser	Ser	Ser
	130					135					140				
Leu	His	Phe	Gly	Asp		Asp	Thr	Val	Thr		Asp	Glu	Asp	Lys	
145					150					155					160
Val	Ser	Val	Arg		Ser	Gln	Thr	He		Asn	Ala	Lys	Ser		Ser
	_			165			_		170					175	
His	Ser	Ala		Ser	His	Lys	Trp		Arg	Thr	Glu	Thr		Ser	Val
			180					185					190		
Ser	Gly		Leu	Met	Lys	Arg		Cys	Leu	li1s	Gly		Ser	Leu	Arg
	,	195	C		,		200		,			205	c		
Arg		Pro	Cys	Arg	Lys	Arg	Phe	Val	Lys	Asn		Ser	Ser	Gin	Arg
Tha	210	Luc	Cln	1	C1	215	11.	Lau	Mat	C15	220	Luc	Lua	A 22 ~	C1
	GIN	Lys	GIN	Lys	230	Arg	11e	Leu	мет	235	Arg	Lys	Lys	Arg	
225	Lou	110	Ana	Ana		Tun	A10	Lou	Lou		Con.	San	Sor	Sor	240
val	Leu	wia	m'g	245	LyS	Tyr	wia	Leu	250	F.1.O	261.	oer.	Sel.	255	ser
Sor	Glu	Acn	Aen		Sor	Ser	Glu	Sor		Sor	Sor	Ser	Ser		Glu
561	010	11011	260	Leu	001	001	GIU	265	501	JUI	001	001	270	1111	GIU

Gly	Glu	Glu	Asp	Leu	Phe	Val	Ser	Ala	Ser	Glu	Asn	His	Gln	Asn	Asn
		275					280					285			
Pro	Ala	Val	Pro	Ser	Gly	Ser	lle	Asp	Glu	Asp	Val	Val	Val	lle	Glu
	290				·	295					300				
Ala	Ser	Ser	Thr	Pro	Gln	Val	Thr	Ala	Asn	Glu	Glu	lle	Asn	Val	Thr
305					310					315					320
Ser	Thr	Asp	Ser	Glu	Val	G1u	He	Val	Thr	Val	Gly	Glu	Ser	Tyr	Arg
				325					330					335	
Ser	Arg	Ser	Thr	Leu	Gly	His	Ser	Arg	Ser	His	Trp	Ser	Gln	Gly	Ser
			340					345					350		
Ser	Ser	His	Ala	Ser	Arg	Pro	Gln	Glu-	Pro	Arg	Asn	Arg	Ser	Arg	Ile
		355					360					365			
Ser	Thr	Val	He	Gln	Pro	Leu	Arg	Gln	Asn	Ala	Ala	Glu	Val	Val	Asp
	370					375					380				
Leu	Thr	Val	Asp	Glu	Asp	Glu	Pro	Thr	Val	Val	Pro	Thr	Thr	Ser	Ala
385					390					395					400
Arg	Met	Glu	Ser	Gln	Ala	Thr	Ser	Ala	Ser	lle	Asn	Asn	Ser	Asn	Pro
				405					410					415	
Ser	Thr	Ser	Glu	Gln	Ala	Ser	Asp	Thr	Ala	Ser	Ala	Val	Thr	Ser	Ser
			420					425					430		
Gln	Pro	Ser	Thr	Val	Ser	Glu	Thr	Ser	Ala	Thr	Leu	Thr	Ser	Asn	Ser
		435					440					445			
Thr	Thr	Gly	Thr	Ser	He	Gly	Asp	Asp	Ser	Arg	Arg	Thr	Thr	Ser	Ser
	450					455					460				
Ala	Val	Thr	Glu	Thr	Gly	Pro	Pro	Ala	Met	Pro	Arg	Leu	Pro	Ser	Cys
465					470					475					480
Cys	Pro	Gln	His	Ser	Pro	Cys	Gly	Gly	Ser	Ser	Gln	Asn	His	His	Ala
				485					490					495	
Leu	Gly	His	Pro	His	Thr	Ser	Cys	Phe	Gln	Gln	His	Gly	His	His	Phe
			500					505					510		
Gln	His	His	His	His	llis	His	His	Thr	Pro	His	Pro	Ala	Val	Pro	Val
		515					520					525			
Ser	Pro	Ser	Phe	Ser	Asp	Pro	Ala	Cys	Pro	Val	Glu	Arg	Pro	Pro	Gln
	530					535					540				
Val	Gln	Ala	Pro	Cys	Gly	Λla	Asn	Ser	Ser	Ser	Gly	Thr	Ser	Tyr	
545					550					555					560

Glu	Gln	Gln	Ala	Leu	Pro	Val	Asp	Leu	Ser	Asn	Ser	Gly	Пlе	Arg	Ser
				565					570					575	
His	Gly	Ser	Gly	Ser	Phe	His	G]y	Ala	Ser	Ala	Phe	Asp	Pro	Cys	Cys
			580					585					590		
Pro	Val	Ser	Ser	Ser	Arg	Ala	Ala	He	Phe	Gly	His	Gln	Ala	Ala	Λla
		595					600					605			
Ala	Ala	Pro	Ser	Gln	Pro	Leu	Ser	Ser	Πe	Asp	Gly	Tyr	Gly	Ser	Ser
	610					615					620				
Met	Val	Ala	Gln	Pro	Gln	Pro	Gln	Pro	Pro	Pro	Gln	Pro	Ser	Leu	Ser
625					630					635					640
Ser	Cys	Arg	His	Tyr	Met	Pro	Pro	Pro	Tyr	Ala	Ser	Leu	Thr	Arg	Pro
				645					650					655	
Leu	His	His	Gln	Ala	Ser	Ala	Cys	Pro	His	Ser	His	Gly	Asn	Pro	Pro
			660					665					670		
Pro	Gln	Thr	Gln	Pro	Pro	Pro	Gln	Val	Asp	Tyr	Val	lle	Pro	His	Pro
		675					680					685			
Val	His	Ala	Phe	His	Ser	Gln	He	Ser	Ser	His	Ala	Thr	Ser	His	Pro
	690					695					700				
Val	Ala	Pro	Pro	Pro	Pro	Thr	His	Leu	Ala	Ser	Thr	Ala	Ala	Pro	lle
705					710					715					720
Pro	Gln	His	Leu	Pro	Pro	Thr	His	Gln	Pro	He	Ser	His	His	lle	Pro
				725					730					735	
Ala	Thr	Ala	Pro	Pro	Ala	GIn	Arg	Leu	His	Pro	His	Glu	Val	Met	Gln
			740					745					750		
Arg	Met	Glu	Va]	Gln	Arg	Arg	Arg	Met	Met	Gln	His	Pro	Thr	Arg	Ala
		755					760					765			
His		Arg	Pro	Pro	Pro	His	Pro	His	Arg	Met		Pro	Asn	Tyr	Gly
	770					775					780			_	
	Gly	His	llis	lle		Val	Pro	Gln	Thr		Ser	Ser	His	Pro	
785					790		_			795					800
G1n	Ala	Pro	Glu		Ser	Ala	Trp	Glu		Gly	Лe	Glu	Ala		Val
				805		•			810		**			815	
Thr	Ala	Ala		Tyr	Thr	Pro	Gly		Leu	His	Pro	His		Ala	His
T		4.3	820	13				825	,	C1		C1	830	,	13
lyr	HIS	Ala	Pro	Pro	Arg	Leu	H1S	H1S	Leu	Ыn	Leu	Gly eas	Ala	Leu	rro

Leu Met Val Pro Asp Met Ala Gly Tyr Pro His Ile Arg Tyr Ile Ser 855 Ser Gly Leu Asp Gly Thr Ser Phe Arg Gly Pro Phe Arg Gly Asn Phe 865 870 875 880 Glu Glu Leu Ile His Leu Glu Glu Arg Leu Gly Asn Val Asn Arg Gly 890 Ala Ser Gln Gly Thr 11e Glu Arg Cys Thr Tyr Pro His Lys Tyr Lys 900 905 Lys Val Thr Thr Asp Trp Phe Ser Gln Arg Lys Leu His Cys Lys Gln 915 920 925 Asp Gly Glu Glu Gly Thr Glu Glu Asp Thr Glu Glu Lys Cys Thr Ile 935 940 Cys Leu Ser Ile Leu Glu Glu Gly Glu Asp Val Arg Arg Leu Pro Cys 945 950 955 960 Met His Leu Phe His Gln Val Cys Val Asp Gln Trp Leu Val Thr Asn 965 970 Lys Lys Cys Pro Ile Cys Arg Val Asp Ile Glu Ala Gln Leu Pro Ser 985 990 Glu Ser

<210> 4825

<211> 105

<212> PRT

<213> Homo sapiens

<400> 448

 Met Glu Gln Ser Trp
 Trp Thr Glu Asn Asp Phe Asp Glu Leu Arg Glu Glu

 1
 5
 10
 15

 Gly Phe Arg Arg Ser Asn Tyr Glu Leu Gln Glu Glu Glu Ile Gln Ile Lys
 20
 25
 30

 Gly Lys Glu Val Lys Asn Phe Glu Lys Asn Leu Asp Glu Cys Ile Thr
 35
 40
 45

 Arg 11e Thr Asn Thr Glu Lys Cys Leu Glu Asp Leu Met Glu Leu Lys

50 55 60

Ala Lys Ala Arg Glu Leu Cys Glu Glu Cys Arg Ser Leu Arg Ser Arg
65 70 75 80

Cys Asp Gln Leu Glu Glu Arg Val Ser Val Met Glu Asp Glu Met Asn
85 90 95

Glu Met Lys Gln Glu Gly Lys Phe Arg
100 105

<210> 4826

<211> 136

<212> PRT

<213> Homo sapiens

<400> 449

Met Trp Leu Glu Val Arg Ala Leu Leu Leu Ser Cys Val Arg Val Phe
1 5 10 15

Gln His Gly Thr Asp Gly Ser Ser Gly His Ser Leu Ser Gln Gly Leu 20 25 30

Ala Arg Leu Arg His Asp Gly Pro Ser Pro Ala Leu Thr Thr Thr Lys
35 40 45

Ser Gly Arg Val Glu Phe Ala Val Asn Ser Ala Val Pro Met Glu Thr 50 55 60

Cys Val Val Ser Glu Gln Gln Tyr Leu Leu Glu Leu Val Gln Thr Pro 65 70 75 80

Arg Leu Pro Ser Gly Thr Thr Trp Ser Thr Ser Pro Arg Glu Ala Ala 85 90 95

Ala Leu Arg Gly Cys Pro Ala Gln Trp Pro Leu Ser Arg Val Ser Leu 100 105 110

Gln Glu Ala Ser Glu Gln Ser Leu Lys Pro Gly Met Ala Pro Ser Pro 115 120 125

Cys Cys His His Cys Val Leu Thr 130 135

<210> 4827

<211> 250

<212> PRT <213> Homo sapiens <400> 450 Met Ala Glu Gly Gly Ala Ser Lys Gly Gly Gly Glu Glu Pro Gly Lys Leu Pro Glu Pro Ala Glu Glu Glu Ser Gln Val Leu Arg Gly Thr Gly His Cys Lys Trp Phe Asn Val Arg Met Gly Phe Gly Phe Ile Ser Met Ile Asn Arg Glu Gly Ser Pro Leu Asp Ile Pro Val Asp Val Phe Val His Gln Ser Lys Leu Phe Met Glu Gly Phe Arg Ser Leu Lys Glu Gly Glu Pro Val Glu Phe Thr Phe Lys Lys Ser Ser Lys Gly Leu Glu Ser Ile Arg Val Thr Gly Pro Gly Gly Ser Pro Cys Leu Gly Ser Glu Arg Arg Pro Lys Gly Lys Thr Leu Gln Lys Arg Lys Pro Lys Gly Asp Arg Cys Tyr Asn Cys Gly Gly Leu Asp His His Ala Lys Glu Cys Ser Leu Pro Pro Gln Pro Lys Lys Cys His Tyr Cys Gln Ser Ile Met His Met Val Ala Asn Cys Pro His Lys Asn Val Ala Gln Pro Pro Ala Ser Ser Gln Gly Arg Gln Glu Ala Glu Ser Gln Pro Cys Thr Ser Thr Leu Pro Arg Glu Val Gly Gly Gly His Gly Cys Thr Ser Pro Pro Phe Pro Gln Glu Ala Arg Ala Glu Ile Ser Glu Arg Ser Gly Arg Ser Pro Gln Glu

Gly Pro Ser Val Gln Lys Arg Lys Lys Thr 245 250

Ala Ser Ser Thr Lys Ser Ser Ile Ala Pro Glu Glu Gln Ser Lys Lys

```
<211> 748
<212> PRT
⟨213⟩ Homo sapiens
<400> 451
Met Lys Gln Met His Glu Trp Asn Phe Thr Ala Ser Ser Ile Lys Gly
                                     10
 1
Ile Ser Leu Ser Lys Phe Asp Glu Arg Cys Cys Phe Leu Tyr Val His
                                 25
Asp Asn Ser Asp Asp Phe Gln 11e Tyr Phe Ser Thr Glu Glu Gln Cys
         35
                                                  45
                             40
Ser Arg Phe Phe Ser Leu Val Lys Glu Met lle Thr Asn Thr Ala Gly
                                              60
                         55
Ser Thr Val Glu Leu Glu Gly Glu Thr Asp Gly Asp Thr Leu Glu Tyr
                                         75
                     70
Glu Tyr Asp His Asp Ala Asn Gly Glu Arg Val Val Leu Gly Lys Gly
Thr Tyr Gly lle Val Tyr Ala Gly Arg Asp Leu Ser Asn Gln Val Arg
                                105
lle Ala Ile Lys Glu Ile Pro Glu Arg Asp Ser Arg Tyr Ser Gln Pro
                                                 125
                            120
Leu His Glu Glu Ile Ala Leu His Lys Tyr Leu Lys His Arg Asn Ile
    130
                        135
Val Gln Tyr Leu Gly Ser Val Ser Glu Asn Gly Tyr Ile Lys Ile Phe
                                        155
                    150
Met Glu Gln Val Pro Gly Gly Ser Leu Ser Ala Leu Leu Arg Ser Lys
                165
                                     170
Trp Gly Pro Met Lys Glu Pro Thr Ile Lys Phe Tyr Thr Lys Gln Ile
            180
                                185
Leu Glu Gly Leu Lys Tyr Leu His Glu Asn Gln lle Val His Arg Asp
                             200
                                                 205
        195
lle Lys Gly Asp Asn Val Leu Val Asn Thr Tyr Ser Gly Val Val Lys
```

215

220

<210> 4828

210

He	Ser	Asp	Phe	Gly	Thr	Ser	Lys	Arg	Leu	Ala	G1 y	Val	Asn	Pro	Cys
225					230					235					240
Thr	Glu	Thr	Phe	Thr	Gly	Thr	Leu	Gln	Tyr	Met	Ala	Pro	$Gl\mathbf{u}$	He	He
				245					250					255	
Asp	Gln	Gly	Pro	Arg	Gly	Tyr	Gly	Ala	Pro	Ala	Asp	lle	Trp	Ser	Leu
			260					265					270		
Gly	Cys	Thr	He	Ile	Glu	Met	Ala	Thr	Ser	Lys	Pro	Pro	Phe	His	Glu
		275					280					285			
Leu	Gly	Glu	Pro	Gln	Ala	Ala	Met	Phe	Lys	Val	Gly	Met	Phe	Lys	Ile
	290					295					300				
His	Pro	Glu	Ile	Pro	Glu	Ala	Leu	Ser	Ala	Glu	Ala	Arg	Ala	Phe	He
305					310					315					320
Leu	Ser	Cys	Phe	Glu	Pro	Asp	Pro	His	Lys	Arg	Ala	Thr	Thr	Ala	Glu
				325					330					335	
Leu	Leu	Arg	Glu	Gly	Phe	Leu	Arg	Gln	Val	Asn	Lys	Gly	Lys	Lys	Asn
			340					345					350		
Arg	Ile	Ala	Phe	Lys	Pro	Ser	Glu	Gly	Pro	Arg	Gly	Val	Val	Leu	Ala
		355					360					365			
Leu	Pro	Thr	Gln	Gly	Glu	Pro	Met	Ala	Thr	Ser	Ser	Ser	Glu	His	Gly
	370					375					380				
Ser	Val	Ser	Pro	Asp	Ser	Asp	Ala	Gln	Pro	Asp	Ala	Leu	Phe	Glu	Arg
385					390					395					400
Thr	Arg	Ala	Pro	Arg	His	His	Leu	Gly	His	Leu	Leu	Ser	Val	Pro	Asp
				405					410					415	
Glu	Ser	Ser	Ala	Leu	Glu	Asp	Arg	Gly	Leu	Ala	Ser	Ser	Pro	Glu	Asp
			420					425					430		
Arg	Asp		Gly	Leu	Phe	Leu		Arg	Lys	Asp	Ser		Arg	Arg	Ala
		435					440					445			
He		Tyr	Lys	Ile	Leu		Glu	Glu	Gln	Asn		Va]	Ala	Ser	Asn
	450					455					460				
	Gln	Glu	Cys	Val		Gln	Ser	Ser	Glu		Leu	His	Leu	Ser	
465					470				_	475					480
Gly	His	He	Lys		He	He	GIy	He		Arg	Asp	Phe	He		Ser
	a :			485			æ,	an)	490	6	-		,	495	
Pro	Glu	His		Val	Met	Ala	Thr		He	Ser	Lys	Leu	Lys	Val	Asp
			500					505					510		

Leu	Asp	Phe	Asp	Ser	Ser	Ser	He	Ser	Gln	Ile	His	Leu	Val	Leu	Phe
		515					520					525			
Gly	Phe	Gln	Asp	Ala	Val	Asn	Lys	Ile	Leu	Arg	Asn	His	Leu	lle	Arg
	530					535					540				
Pro	His	Trp	Met	Phe	Ala	Met	Asp	Asn	He	He	Arg	Arg	Ala	Val	Gln
545					550					555					560
Ala	Ala	Val	Thr	11e	Leu	Ile	Pro	Glu	Leu	Arg	Ala	His	Phe	Glu	Pro
				565					570					575	
Thr	Cys	Glu	Thr	Glu	Gly	Val	Asp	Lys	Asp	Met	Asp	Glu	Ala	Glu	Glu
			580					585					590		
Gly	Tyr	Pro	Pro	Ala	Thr	Gly	Pro	Gly	Gln	Glu	Ala	Gln	Pro	His	Gln
		595					600					605			
Gln	His	Leu	Ser	Leu	Gln	Leu	•Gly	Glu	Leu	Arg	Gln	Glu	Thr	Asn	Arg
	610					615					620				
Leu	Leu	Glu	His	Leu	Val	Glu	Lys	Glu	Arg	Glu	Tyr	Gln	Asn	Leu	Leu
625					630					635					640
Arg	Gln	Thr	Leu	Glu	Gln	Lys	Thr	Gln	${\sf Glu}$	Leu	Tyr	His	Leu	Gln	Leu
				645					650					655	
Lys	Leu	Lys	Ser	Asn	Cys	Ile	Thr	Glu	Asn	Pro	Ala	Gly	Pro	Tyr	Gly
			660					665					670		
Gln	Arg	Thr	Asp	Lys	Glu	Leu	Ile	Asp	Trp	Leu	Arg	Leu	Gln	Gly	Ala
		675					680					685			
Asp	Ala	Lys	Thr	lle	Glu	Lys	He	Val	Glu	Glu	Gly	Tyr	Thr	Leu	Ser
	690					695					700				
Asp	lle	Leu	Asn	Glu	He	Thr	Lys	Glu	Asp	Leu	Arg	Tyr	Leu	Arg	Leu
705					710					715					720
Arg	Gly	Gly	Leu	Leu	Cys	Arg	Leu	Trp	Ser	Ala	Val	Ser	Gln	Tyr	Arg
				725					730					735	
Arg	Ala	G1n	Glu	Ala	Ser	Glu	Thr	Lys	Asp	Lys	Ala				
			740					745							

<210> 4829

<211> 572

<212> PRT

<213> Homo sapiens

<400)> 45	52													
Met	Cys	His	Phe	Lys	Leu	Va]	Ala	lle	Va]	Gly	Tyr	Leu	lle	Arg	Leu
1				5					10					15	
Ser	lle	Lys	Ser	Ile	Gln	lle	Glu	Ala	Asp	Asn	Cys	Val	Thr	Asp	Ser
			20					25					30		
Leu	Thr	Ile	Tyr	Asp	Ser	Leu	Leu	Pro	lle	Arg	Ser	Ser	He	Leu	Tyr
		35					40					45			
Arg	Ile	Cys	Glu	Pro	Thr	Arg	Thr	Leu	Met	Ser	Phe	Val	Ser	Thr	Asn
	50					55					60				
Asn	Leu	Met	Leu	Val	Thr	Phe	Lys	Ser	Pro	His	Ile	Arg	Arg	Leu	Ser
65					70					75					80
Gly	He	Arg	Ala	Tyr	Phe	Glu	Val	Ile	Pro	Glu	Gln	Lys	Cys	G]u	Asn
				85					90					95	
Thr	Val	Leu	Val	Lys	Asp	Ile	Thr	Gly	Phe	Glu	Gly	Lys	lle	Ser	Ser
			100					105					110		
Pro	Tyr	Tyr	Pro	Ser	Tyr	Tyr	Pro	Pro	Lys	Cys	Lys	Cys	Thr	Trp	Lys
		115					120					125			
Phe	Gln	Thr	Ser	Leu	Ser	Thr	Leu	Gly	Ile	Ala	Leu	Lys	Phe	Tyr	Asn
	130					135					140				
Tyr	Ser	Ile	Thr	Lys	Lys	Ser	Met	Lys	Gly	Cys	Glu	His	Gly	Trp	Trp
145					150					155					160
Glu	lle	Asn	Glu	His	Met	Tyr	Cys	Gly	Ser	Tyr	Met	Asp	His	Gln	Thr
				165					170					175	
He	Phe	Arg	Val	Pro	Ser	Pro	Leu	Val	His	He	Gln	Leu	Gln	Cys	Ser
			180					185					190		
Ser	Arg	Leu	Ser	Asp	Lys	Pro	Leu	Leu	Ala	Glu	Tyr	G1 y	Ser	Tyr	Asn
		195					200					205			
He		Gln	Pro	Cys	Pro		Gly	Ser	Phe	Arg	Cys	Ser	Ser	Gly	Leu
	210					215	_				220		_		
	Val	Pro	Gln	Ala		Arg	Cys	Asp	Gly		Asn	Asp	Cys	Phe	
225			0.1		230			•		235	_				240
Glu	Ser	Asp	Glu		Phe	Cys	Val	Ser		GIn	Pro	Ala	Cys		Thr
				245					250				ъ.	255	
Ser	Ser	Phe		GIn	His	Gly	Pro		Пe	Cys	Asp	Gly		Arg	Asp
			260					265					270		

Cys	Glu	Asn	Gly	Arg	Asp	Glu	Gln	Asn	Cys	Thr	Gln	Ser	Ile	Pro	Cys
		275					280					285			
Asn	Asn	Arg	Thr	Phe	Lys	Cys	Gly	Asn	Asp	lle	Cys	Phe	Arg	Lys	Gln
	290					295					300				
Asn	Ala	Lys	Cys	Asp	Gly	Thr	Val	Asp	Cys	Pro	Asp	Gly	Ser	Asp	Glu
305					310					315					320
Glu	Gly	Cys	Thr	Cys	Ser	Arg	Ser	Ser	Ser	Ala	Leu	His	Arg	11e	lle
				325					330					335	
Gly	Gly	Thr	Asp	Thr	Leu	Glu	Gly	Gly	Trp	Pro	Trp	Gln	Val	Ser	Leu
			340					345					350		
His	Phe	Val	Gly	Ser	Ala	Tyr	Cys	Gly	Ala	Ser	Val	Ile	Ser	Arg	Glu
		355					360					365			
Trp	Leu	Leu	Ser	Ala	Ala	His	Cys	Phe	His	Gly	Asn	Arg	Leu	Ser	Asp
	370					375					380				
Pro	Thr	Pro	Trp	Thr	Ala	His	Leu	Gly	Met	Tyr	Val	Gln	Gly	Asn	Ala
385					390					395					400
Lys	Phe	Val	Ser	Pro	Val	Arg	Arg	He	Val	Val	His	Glu	Tyr	Tyr	Asn
				405					410					415	
Ser	Gln	Thr	Phe	Asp	Tyr	Asp	Ile	Ala	Leu	Leu	Gln	Leu	Ser	lle	Ala
			420					425					430		
Trp	Pro	Glu	Thr	Leu	Lys	Gln	Leu	He	Gln	Pro	Ile	Cys	lle	Pro	Pro
		435					440					445			
Thr	Gly	Gln	Arg	Val	Arg	Ser	Gly	Glu	Lys	Cys	Trp	Val	Thr	Gly	Trp
	450					455					460				
Gly	Arg	Arg	His	Glu	Ala	Asp	Asn	Lys	Gly	Ser	Leu	Val	Leu	G1n	Gln
465					470					475	•				480
Ala	Glu	Va]	Glu	Leu	lle	Asp	G1n	Thr	Leu	Cys	Val	Ser	Thr	Tyr	Gly
				485					490					495	
He	He	Thr	Ser	Arg	Met	Leu	Cys	Ala	Gly	Ile	Met	Ser	Gly	Lys	Arg
			500					505					510		
Asp	Ala	Cys	Lys	Gly	Asp	Ser	Gly	Gly	Pro	Leu	Ser	Cys	Arg	۸rg	Lys
		515					520					525			
Ser	Asp	Gly	Lys	Trp	lle	Leu	Thr	Gly	He	Val	Ser	Trp	Gly	His	Gly
	530					535					540				
Cys	Gly	Arg	Pro	Asn	Phe	Pro	Gly	Val	Tyr	Thr	Arg	Val	Ser	Asn	Phe
545					550					555					560

Val Pro Trp Ile His Lys Tyr Val Pro Ser Leu Leu 565 570

<210> 4830

<211> 134

<212> PRT

<213> Homo sapiens

<400> 453

Met Val Gly Leu Leu Phe His Ala Pro Lys Ala Pro Glu Met Ala Pro 1 5 10 15

Leu Arg Cys Cys lle Met Asn Lys Ile Ile Met Val Arg Arg Pro Lys
20 25 30

Gln Ser Thr Ala Asp Tyr Gly Met Arg Thr Ser Gly Pro Val Glu Ser 35 40 45

Gly Leu Ser Ala Asp Ser Leu Gln Leu Leu Cys Ser Tyr Ala Ala 11e 50 55 60

Lys Asn Ser Ala Glu Leu Leu Met Val Gly Pro Gln Gly Met Arg Pro 65 70 75 80

Ala Thr Gly Gln Asp Leu Leu Cys Arg Pro Cys Leu Ser His Asp Leu 85 90 95

Pro Gly Pro Leu His Pro Pro Arg Gly Leu Ser Gly Ser Ser Ser Leu
100 105 110

Leu Ile Ser Pro Arg Leu Gln Asp Val Ser Leu Gln Leu Val His Pro 115 120 125

Thr Pro Glu Glu Ser Phe 130

<210> 4831

<211> 184

<212> PRT

<213> Homo sapiens

<400> 454

Met Pro Phe Arg Lys Ala Cys Gly Pro Lys Leu Thr Asn Ser Pro Thr 10 Val Ile Val Met Val Gly Leu Pro Ala Arg Gly Lys Thr Tyr Ile Ser 20 25 30 Lys Lys Leu Thr Arg Tyr Leu Tyr Trp Ile Gly Val Pro Thr Lys Val 40 Phe Asn Val Gly Glu Tyr Arg Arg Glu Ala Val Lys Gln Tyr Ser Ser 55 Tyr Asn Phe Phe Arg Pro Asp Asn Glu Glu Ala Met Lys Val Arg Lys 65 70 Gln Cys Ala Leu Ala Ala Leu Arg Asp Val Lys Ser Tyr Leu Ala Lys 90 Glu Gly Gly Gln 11e Ala Val Phe Asp Ala Thr Asn Thr Thr Arg Glu 100 105 110 Arg Arg His Met Ile Leu His Phe Ala Lys Glu Asn Asp Phe Lys Ala 125 120 Phe Phe Ile Glu Ser Val Cys Asp Asp Pro Thr Val Val Ala Ser Asn 135 lle Met Gln Lys Ala Phe Gln Arg Asp Leu Ser Ala Thr Pro Leu Trp 145 155 160 Cys Cys Arg Lys Leu Lys Ser Pro Ala Arg Ile Thr Lys Thr Ala Thr 165 170 175 Arg Gln Lys Pro Trp Thr Thr Ser 180

<210> 4832

<211> 493

<212> PRT

<213> Homo sapiens

<400> 455

Met Gly Pro Thr Ser Val Leu Arg Ala Gly Leu Thr Pro Ser Cys Leu

1 5 10 15

Pro Pro Pro Pro Ser Gly Ala Thr Asn Gly Ser Val Ser Pro Leu Gly Arg
20 25 30

Ala	Gln	Arg	Val	Trp	Thr	Glu	Pro	Gly	Gly	Arg	Gly	Leu	His	Gly	Ala
		35					40					45			
Thr	Ala	Ala	Gly	Pro	Val	Ala	Ala	Ala	Cys	Pro	Leu	Leu	Ala	Val	Thr
	50					55					60				
Ala	Thr	Ala	Pro	Gly	Gln	Pro	Ser	Gly	Ala	Ser	Thr	Val	Trp	Val	Arg
65					70					75					80
Glu	Gly	Gly	Thr	Ala	Pro	Ala	Thr	Arg	Met	Thr	Val	Pro	Leu	Ala	Pro
				85					90					95	
Arg	Thr	Ser	Glu	Lys	Cys	Ser	Val	Leu	Asn	Leu	Thr	Ala	Ser	Leu	Ser
			100					105					110		
Val	Gly	Asn	Ser	Thr	Ser	Gly	Lys	Arg	Thr	Gly	Glu		Cys	Ser	Leu
		115					120					125			
Thr	Cys	Leu	Ala	Glu	Gly		Asn	Phe	Tyr	Thr	Glu	Arg	Ala	Ala	Ala
	130					135					140				
Val	Val	Asp	Gly	Thr	Pro	Cys	Arg	Pro	Asp		Val	Asp	Ile	Cys	
145					150					155					160
Ser	Gly	Glu	Cys		His	Val	Gly	Cys		Arg	Val	Leu	Gly		Asp
				165					170					175	
Leu	Arg	Glu		Lys	Cys	Arg	Val		Gly	Gly	Asp	Gly		Ala	Cys
			180			ъ.		185				0.1	190		
Glu	Thr		Glu	GIy	Val	Phe		Pro	Ala	Ser	Pro		Ala	Gly	Tyr
0.1		195				Б	200	0.1				205	D.	* *	0.1
Glu		val	Val	Trp	lle		Lys	Gly	Ser	Val		11e	Phe	He	GIn
	210			C	,	215		•	4.7	,	220	61		C1	C1
	Leu	Asn	Leu	Ser	Leu	Asn	HIS	Leu	Ala		Lys	GIY	Asp	GIN	
225	Lau	Lau	Lau	C1	230	Lau	Duo	C1	Thu	235 Pro	Cln	Dwa	uio	Ana	240
ser	Leu	Leu	Leu		Gly	Leu	F10	GTy		110	GIN	F10	шѕ	255	Leu
Dreo	Lou	A10	Cly	245	Thr	Dho	Cln	Lou	250	Cln	Cly	Dro	Acn		Va 1
110	Leu	МІа	260	1111	1111	1116	UIII	265	MI g	GIII	O1 y	110	270	OIII	vai
Gln	Sor	Lou		Δ1a	Leu	C1v	Pro		Acn	412	Sor	Lou		Val	Mot
0111	361	275	Ulu	пта	Leu	Gry	280	116	ASII	пта	261	285	116	vai	мес
Val	Lou		Ara	Thr	Glu	Lou		Ala	Lau	Ara	Tur		Pho	Aen	Λla
, ci 1	290	ma	mg	1111	010	295	110	11.1.0	LCU	мg	300	m g	1116	11-511	HIG
Pro		Ala	Arø	Asp	Ser		Pro	Pro	Tvr	Ser		His	Tvr	Ala	Pro
305		,110	6	Ф	310	200			. , .	315		,,15	.,.	1110	320

Trp Thr Arg Cys Ser Ala Gln Cys Ala Gly Gly Ser Gln Val Gln Ala Val Glu Cys Arg Asn Gln Leu Asp Gly Ser Ala Val Ala Pro His Tyr Cys Ser Ala His Ser Lys Leu Pro Lys Arg Gln Arg Ala Cys Asn Thr Glu Pro Cys Pro Pro Asp Trp Val Val Gly Asn Trp Ser Leu Cys Ser Arg Ser Cys Asp Ala Gly Val Arg Ser Arg Ser Val Val Cys Gln Arg Arg Val Ser Ala Ala Glu Glu Lys Ala Leu Asp Asp Ser Ala Cys Pro Gln Pro Arg Pro Pro Val Leu Glu Ala Cys His Gly Pro Thr Cys Pro Pro Glu Trp Ala Ala Leu Asp Trp Ser Glu Cys Thr Pro Ser Cys Gly Pro Gly Leu Arg His Arg Val Val Leu Cys Lys Ser Ala Asp His Arg Ala Thr Leu Pro Pro Ala His Cys Ser Pro Ala Ala Lys Pro Pro Ala Thr Met Arg Cys Asn Leu Arg Arg Cys Pro Pro Ala Arg

<210> 4833

<211> 803

<212> PRT

<213> Homo sapiens

<400> 456

 Met 11e Ser Ala Gln Cys Asn Leu Cys Leu Pro Gly Ser Gly Asp Ser

 1
 5
 10
 15

 Cys Ala Ser Gly Ser Gln Leu Ala Gly Thr Thr Val Ala Gly Pro Ala
 20
 25
 30

 His Pro Glu Ser Leu Thr Leu Cys Ser Gly Glu Ala Glu Arg Arg Gly

35 40 45

Lys	Glu	Glu	Trp	Pro	Asp	Ser	His	His	Tyr	Pro	G1 y	Pro	Thr	Pro	Ala
	50					55					60				
Pro	Glu	Ser	Ser	Asp	Gly	Pro	His	Lys	Val	Thr	Val	Leu	Ala	Thr	Met
65					70					75					80
Leu	Ser	Ser	Arg	Trp	Trp	Pro	Ser	Ser	Trp	Gly	He	Leu	Gly	Leu	Gly
				85	•				90					95	
Pro	Arg	Ser	Pro	Pro	Arg	Gly	Ser	Gln	Leu	Cys	Λla	Leu	Tyr	Ala	Phe
			100					105					110		
Thr	Tyr	Thr	Gly	Ala	Asp	Gly	Gln	Gln	Val	Ser	Leu	Ala	Glu	Gly	Asp
		115					120					125			
Arg	Phe	Leu	Leu	Leu	Arg	Lys	Thr	Asn	Ser	Asp	Trp	Trp	Leu	Ala	Arg
	130					135					140				
Arg	Leu	Glu	Ala	Pro	Ser	Thr	Ser	Arg	Pro	He	Phe	Val	Pro	Ala	Ala
145					150					155					160
Tyr	Met	lle	Glu	Glu	Ser	He	Pro	Ser	Gln	Ser	Pro	Thr	Thr	Val	He
				165					170					175	
Pro	Gly	Gln	Leu	Leu	Trp	Thr	Pro	Gly	Pro	Lys	Leu	Phe	His	Gly	Ser
			180					185					190		
Leu	Glu	Glu	Leu	Ser	Gln	Ala	Leu	Pro	Ser	Arg	Ala	Gln	Ala	Ser	Ser
		195					200					205			
Glu	Gln	Pro	Pro	Pro	Leu	Pro	Arg	Lys	Met	Cys	Arg	Ser	Val	Ser	Thr
	210					215					220				
Asp	Asn	Leu	Ser	Pro	Ser	Leu	Leu	Lys	Pro	Phe	Gln	Glu	Gly	Pro	Ser
225					230					235					240
Gly	Arg	Ser	Leu	Ser	Gln	Glu	Asn	Leu	Pro	Pro	Glu	Ala	Ser	Ala	Ser
				245					250					255	
Thr	Ala	Gly	Pro	Gln	Pro	Leu	Met	Ser	Glu	Pro	Pro	Val	Tyr	Cys	Asn
			260					265					270		
Leu	Val	Asp	Leu	Arg	Arg	Cys	Pro	Arg	Ser	Pro	Pro	Pro	G1 y	Pro	Ala
		275					280					285			
Cys	Pro	Leu	Leu	Gln	Arg	Pro	Asp	Ala	Trp	Glu	Gln	His	Leu	Asp	Pro
	290					295					300				
Asn	Ser	Gly	Arg	Cys	Phe	Tyr	lle	Asn	Ser	Leu	Thr	Gly	Cys	Lys	Ser
305					310					315					320
Trp	Lys	Pro	Pro	Arg	Arg	Ser	Arg	Ser	Glu	Thr	Asn	Pro	Gly	Ser	Met
				325					330					335	

Glu	Gly	Thr	Gln	Thr	Leu	Lys	Arg	Asn	Asn	Asp	Val	Leu	Gln	Pro	Gln
			340					345					350		
Ala	Lys	Gly	Phe	Arg	Ser	Asp	Thr	Gly	Thr	Pro	Glu	Pro	Leu	Asp	Pro
		355					360					365			
Gln	Gly	Ser	Leu	Ser	Leu	Ser	Gln	Arg	Thr	Ser	Gln	Leu	Asp	Pro	Pro
	370					375					380				
Λla	Leu	Gln	Ala	Pro	Arg	Pro	Leu	Pro	Gln	Leu	Leu	Asp	Asp	Pro	His
385					390					395					400
Glu	Val	Glu	Lys	Ser	Gly	Leu	Leu	Asn	Met	Thr	Lys	lle	Ala	Gln	Gly
				405					410					415	
Gly	Arg	Lys	Leu	Arg	Lys	Asn	Trp	Gly	Pro	Ser	Trp	Val	Val	Leu	Thr
			420					425					430		
Gly	Asn	Ser	Leu	Val	Phe	Tyr	Arg	Glu	Pro	Pro	Pro	Thr	Ala	Pro	Ser
		435					440					445			
Ala	Gly	Trp	Gly	Pro	Ala		Ser	Arg	Pro	Glu	Ser	Ser	Val	Asp	Leu
	450					455					460				
Arg	Gly	Ala	Ala	Leu	Ala	His	Gly	Arg	His		Ser	Ser	Arg	Arg	
465					470					475					480
Val	Leu	His	lle		Thr	lle	Pro	Gl y		Glu	Phe	Leu	Leu		Ser
				485				_	490					495	
Asp	His	Glu		Glu	Leu	Arg	Ala		His	Arg	Ala	Leu		Thr	Val
			500					505					510		- 1
He	Glu		Leu	Asp	Arg	Glu		Pro	Leu	Glu	Leu		Leu	Ser	Gly
	0.7	515	. 7	6.7			520	0.1	0.1		0.1	525	0.1	0.1	~
Ser		Pro	Ala	Glu	Leu		Ala	Gly	Glu	Asp		Glu	Glu	Glu	Ser
<i>a</i> 1	530		C	,	D	535	,			C	540				0
	Leu	val	Ser	Lys	Pro	Leu	Leu	Arg	Leu		Ser	Arg	Arg	Ser	
545	Α	C1	D	C1	550	The	C1	C1	Α	555	v 1	A	Δ	1	560
116	Arg	GIŸ	Pro		Gly	Inr	GIU	GIN		Arg	vai	Arg	Asn		Leu
1	Δ	1	11.	565	1	A	Dana	Dua	570	C1	C	Lan	C1	575	A
Lys	Arg	Leu		Ala	Lys	Arg	Pro	585	Leu	GIN	ser	Leu	590	61 u	Arg
C1	Lau	Lau	580	Aan	Cla	Vol	Dho		Cvc	Cln	Lan	Clu		Lau	Cva
ОТУ	Leu		Arg	иер	Gln	var		GIŸ	Cys	GIII	Leu	605	Ş€1	Leu	Cys
61.	Arc	595	G1v	Acr	Thr	Val	600 Pro	Son	Pho	Low	A 25.00		Cvc	11.	A16
0111	610	OIU	ΩIŅ	nsp	141.	615	110	Set.	1116	Leu	Arg 620	Leu	Cys	116	ита
	010					010					UZU				

Ala Val Asp Lys Arg Gly Leu Asp Val Asp Gly Ile Tyr Arg Val Ser 625 630 Gly Asn Leu Ala Val Val Gln Lys Leu Arg Phe Leu Val Asp Arg Glu 650 Arg Ala Val Thr Ser Asp Gly Arg Tyr Val Phe Pro Glu Gln Pro Gly 665 Gln Glu Gly Arg Leu Asp Leu Asp Ser Thr Glu Trp Asp Asp 11e His 680 685 Val Val Thr Gly Ala Leu Lys Leu Phe Leu Arg Glu Leu Pro Gln Pro 700 690 695 Leu Val Pro Pro Leu Leu Leu Pro His Phe Arg Ala Ala Leu Ala Leu 710 715 Ser Gln 11e Gln Glu Leu 11e Gly Ser Met Pro Lys Pro Asn His Asp 730 725 735 Thr Leu Arg Tyr Leu Leu Glu His Leu Cys Arg Val Ile Ala His Ser 745 Asp Lys Asn Arg Met Thr Pro His Asn Leu Gly 11e Val Phe Gly Pro 760 Thr Leu Phe Arg Pro Glu Gln Glu Thr Ser Asp Pro Ala Ala His Ala 770 775 780 Leu Tyr Pro Gly Gln Leu Val Gln Leu Met Leu Thr Asn Phe Thr Ser 790 795 800 785 Leu Phe Pro

<210> 4834

<211> 892

<212> PRT

<213> Homo sapiens

<400> 457

 Met Gln Asp Leu Lys Lys Lys Tyr Lys Lys Lys Lys Gln Lys Arg Met Lys

 1
 5
 10
 15

 Ser Arg Lys Val Arg Lys Pro Thr Glu Asn Gln Glu Lys Asn Ile Arg
 20
 25
 30

Lys	Arg	Glu 35	Arg	Arg	Lys	Lys	Ser 40	Lys	Arg	Arg	Lys	Arg 45	Glu	Lys	His
Lys	His 50	Asn	Ser	Pro	Ser	Ser 55	Asp	Asp	Ser	Ser	Asp	Tyr	Ser	Leu	Лsp
Ser 65	Asp	Val	Glu	His	Thr 70	Glu	Ser	Ser	His	Lys 75	Lys	Arg	Thr	Gly	Phe 80
Tyr	Arg	Asp	Tyr	Лsр 85	lle	Pro	Phe	Thr	G1n 90	Arg	Gly	His	lle	Ser 95	Gly
Ser	Tyr	Ile	Thr 100	Ser	Lys	Lys	Gly	Gln 105	His	Λsn	Lys	Lys	Phe 110	Lys	Ser
Lys	Glu	Tyr 115	Asp	Glu	Tyr	Ser	Thr 120	Tyr	Ser	Asp	Asp	Asn 125	Phe	Gly	Asn
Tyr	Ser 130	Asp	Asp	Asn	Phe	Gly 135	Asn	Tyr	Gly	Gln	Glu 140	Thr	Glu	Glu	Asp
Phe 145	Ala	Asn	Gln	Leu	Lys 150	Gln	Tyr	Arg	Gln	Ala 155	Lys	Glu	Thr	Ser	Asn 160
lle	Ala	Leu	Gly	Ser 165	Ser	Phe	Ser	Lys	G] u 170	Ser	Gly	Lys	Lys	Gln 175	Arg
Met	Lys	Gly	Val 180	Gln	Gln	Gly	Ile	Glu 185	Gln	Arg	Val	Lys	Ser 190	Phe	Asn
Val	Gly	Arg 195	Gly	Arg	Gly	Leu	Pro 200	Lys	Lys	lle	Lys	Arg 205	Lys	Glu	Arg
Gly	Gly 210	Arg	Thr	Asn	Lys	Gly 215	Pro	Asn	Val	Phe	Ser 220	Val	Ser	Asp	Asp
Phe 225	Gln	Glu	Tyr	Asn	Lys 230	Pro	Gly	Lys	Lys	Trp 235	Lys	Val	Met	Thr	Gln 240
Glu	Phe	lle	Asn	Gln 245	His	Thr	Val	Glu	His 250	Lys	Gly	Lys	Gln	lle 255	Cys
Lys	Tyr	Phe	Leu 260	Glu	Gly	Arg	Cys	11e 265	Lys	Gly	Asp	Gln	Cys 270	Lys	Phe
Asp	His	Asp 275	Ala	Glu	Leu	Glu	Lys 280	Arg	Lys	Glu	11e	Cys 285	Lys	Phe	Tyr
Leu	Gln 290	Gly	Tyr	Cys	Thr	Lys 295	Gly	Głu	Asn	Cys	11e 300	Tyr	Met	His	Asn
G1u 305	Phe	Pro	Cys	Lys	Phe 310	Tyr	His	Ser	Gly	Ala 315	Lys	Cys	Tyr	Gln	Gly 320

Asp	Asn	Cys	Lys	Phe	Ser	His	Asp	Asp	Leu	Thr	Lys	Glu	Thr	Lys	Lys
				325					330					335	
Leu	Leu	Asp	Lys	Val	Leu	Λsn	Thr	Asp	Glu	Glu	Leu	He	Asn	Glu	Asp
			340					345					350		
Glu	Arg	Glu	Leu	Glu	Glu	Leu	Arg	Lys	Arg	Gly	He	Thr	Pro	Leu	Pro
		355					360					365			
Lys	Pro	Pro	Pro	Gly	Val	Gly	Leu	Leu	Pro	Thr	Pro	Pro	Glu	His	Phe
	370					375					380				
Pro	Phe	Ser	Asp	Pro	Glu	Asp	Asp	Phe	Gln	Thr	Asp	Phe	Ser	Asp	Asp
385					390					395				•	400
Phe	Arg	Lys	Ile	Pro	Ser	Leu	Phe	Glu	He	Val	Val	Lys	Pro	Thr	Val
				405					410					415	
Asp	Leu	Ala	His	Lys	He	G1 y	Arg	Lys	Pro	Pro	Ala	Phe	Tyr	Thr	Ser
			420					425	•				430		
Ala	Ser	Pro	Pro	Gly	Pro	Gln	Phe	Gln	G1 y	Ser	Ser	Pro	His	Pro	Gln
		435					440					445			
His	lle	Tyr	Ser	Ser	Gly	Ser	Ser	Pro	Gly	Pro	Gly	Pro	Asn	Met	Ser
	450					455					460				
Gln	Gly	His	Ser	Ser	Pro	Val	Met	His	Pro	Gly	Ser	Pro	Gly	His	His
465					470					475					480
Pro	Cys	Ala	Gly	Pro	Pro	Gly	Leu	Pro	Val	Pro	Gln	Ser	Pro	Pro	Leu
				485					490					495	
Pro	Pro	Gly	Pro	Pro	Glu	lle	Val	Gly	Pro	Gln	Asn	Gln	Ala	Gly	Val
			500					505					510		
Leu	Val	Gln	Pro	Asp	Thr	Ser	Leu	Thr	Pro	Pro	Ser	Met	Gly	Gly	Ala
		515					520					525			
Tyr	His	Ser	Pro	Gly	Phe	Pro	Gly	His	Val	Met	Lys	Val	Pro	Arg	Glu
	530					535					540				
Asn	His	Cys	Ser	Pro	Gly	Ser	Ser	Tyr	Gln	Gln	Ser	Pro	Gly	Glu	Met
545					550					555					560
Gln	Leu	Asn	Thr	Asn	Tyr	Glu	Ser	Leu	Gln	Asn	Pro	Ala	Glu	Phe	Tyr
				565					570					575	
Asp	Asn	Tyr	Tyr	Ala	Gln	His	Ser	lle	His	Asn	Phe	Gln	Pro	Pro	Asn
			580					585					590		
Asn	Ser	Gly	Asp	Gly	Met	Trp	His	Gly	Glu	Phe	Ala	Gln	Gln	G]n	Pro
		595					600					605			

Pro	Val	Val	Gln	Asp	Ser	Pro	Asn	His	Gly	Ser	Gly	Ser	Asp	G1 y	Ser
	610					615					620				
Ser	Thr	Arg	Thr	Gly	His	Gly	Pro	Leu	Pro	Val	Pro	Gly	Leu	Leu	Pro
625					630					635					640
Ala	Val	Gln	Arg	Ala	Leu	Phe	Val	Arg	Leu	Thr	Gln	Arg	Tyr	Gln	Glu
				645					650					655	
Asp	G1u	Glu	Gln	Thr	Ser	Thr	GIn	Pro	His	Arg	Ala	Pro	Ser	Lys	Glu
			660					665					670		
Glu	Asp	Asp	Thr	Val	Asn	Trp	Tyr	Ser	Ser	Ser	Glu	Glu	Glu	Glu	Gly
		675					680					685			
Ser	Ser	Val	Lys	Ser	lle	Leu	Lys	Thr	Leu	Gln	Lys	Gln	Thr	Glu	Thr
	690					695					700				
Leu	Arg	Asn	Gln	Gln	Gln	Pro	Ser	Thr	Glu	Leu	Ser	Thr	Pro	Thir	Λsp
705					710					715					720
Pro	Arg	Leu	Ala	Lys	Glu	Lys	Ser	Lys	Gly	Asn	Gln	Val	Val	Asp	Pro
				725					730					735	
Arg	Leu	Arg	Thr	lle	Pro	Arg	Gln	Asp	lle	Arg	Lys	Pro	Ser	Glu	Ser
			740					745					750		
Ala	Pro	Leu	Asp	Leu	Arg	Leu	Ala	Trp	Asp	Pro	Arg	Lys	Leu	Arg	Gly
		755					760					765			
Asn	Gly	Ser	Gly	His	lle	Gly	Ser	Ser	Va]	Gly	Gly	Ala	Lys	Phe	Asp
	770					775					780				
Leu	His	His	Ala	Asn	Ala	Gly	Thr	Asn	Val	Lys	His	Lys	Arg	Gly	Asp
785					790					795					800
Asp	Asp	Asp	Glu	Asp	Thr	Glu	Arg	Glu	Leu	G1 y	Glu	Lys	Ala	Phe	Leu
				805					810					815	
lle	Pro	Leu	Asp	Ala	Ser	Pro	Gly	lle	Met	Leu	Gln	Asp	Pro	Arg	Ser
			820					825					830		
Gln	Leu	Arg	Gln	Phe	Ser	His	He	Lys	Lys	Asp	lle	Thr	Leu	Thr	Lys
		835					840					845			
Pro	Asn	Phe	Ala	Lys	His	He	Val	Trp	Ala	Pro	Glu	Asp	Leu	Leu	Pro
	850					855					860				
Val	Pro	Leu	Pro	Lys	Pro	Asp	Pro	Val	Ser	Ser	He	Asn	Leu	Pro	Leu
865					870					875					880
Pro	Pro	Leu	Пe	Ala	Asp	Gln	Arg	Leu	Asn	Arg	Leu				

885 890

<210> 4835

<211> 891 <212> PRT <213> Homo sapiens <400> 458 Met Ala Leu Thr Gln Gly Pro Leu Thr Phe Arg Asp Val Ala lle Glu Phe Ser Gln Glu Glu Trp Lys Ser Leu Asp Pro Val Gln Lys Ala Leu 25 Tyr Trp Asp Val Met Leu Glu Asn Tyr Arg Asn Leu Val Phe Leu Gly 35 40 45 Ile Leu Pro Lys Cys Met Thr Lys Glu Leu Pro Pro Ile Gly Asn Ser 55 60 Asn Thr Gly Glu Lys Cys Gln Thr Val Thr Leu Glu Arg His Glu Cys 70 65 75 80 Tyr Asp Val Glu Asn Phe Tyr Leu Arg Glu Ile Gln Lys Asn Leu Gln 90 Asp Leu Glu Phe Gln Trp Lys Asp Gly Glu lle Asn Tyr Lys Glu Val 105 Pro Met Thr Tyr Lys Asn Asn Leu Asn Gly Lys Arg Gly Gln His Ser · 125 115 120 Gln Glu Asp Val Glu Asn Lys Cys Ile Glu Asn Gln Leu Thr Leu Ser 135 140 Phe Gln Ser Arg Leu Thr Glu Leu Gln Lys Phe Gln Thr Glu Gly Lys 145 150 155 160 Ile Tyr Glu Cys Asn Gln Ser Glu Lys Thr Val Asn Asn Ser Ser Leu 165 170 Val Ser Pro Leu Gln Arg lle Leu Pro Ser Val Gln Thr Asn lle Ser 185 Lys Lys Tyr Glu Asn Glu Phe Leu Gln Leu Ser Leu Pro Thr Gln Leu 200 205 195 Glu Lys Thr His Ile Arg Glu Lys Pro Tyr Ile Cys Lys Gly Cys Gly

	210					215					220				
Lys	Ala	Phe	Arg	Val	Ser	Ser	Ser	Leu	lle	Asn	His	Gln	Met	Val	His
225					230			•		235				•	240
Thr	Thr	Glu	Lys	Pro	Tyr	Lys	Cys	Asn	Glu	Cys	Gly	Lys	Ala	Phe	His
				245					250				-	255	
Arg	Gly	Ser	Leu	Leu	Thr	Ile	His	Gln	Ile	Val	His	Thr	Arg	Gly	Lys
			260					265					270		
Pro	Tyr	Gln	Cys	Gly	Val	Cys	Gly	Lys	He	Phe	Arg	Gln	Asn	Ser	Asp
		275					280					285			
Leu	Val	Asn	His	Arg	Arg	Ser	His	Thr	Gly	Glu	Lys	Pro	Tyr	Lys	Cys
	290					295					300				
Asn	Glu	Cys	Gly	Lys	Ser	Phe	Ser	Gln	Ser	Tyr	Asn	Leu	Ala	He	His
305					310					315					320
Gln	Arg	Ile	His	Thr	Gly	Glu	Lys	Pro	Tyr	Lys	Cys	Asn	Glu	Cys	Gly
				325					330					335	
Lys	Thr	Phe	Lys	Gln	Gly	Ser	Cys	Leu	Thr	Thr	His	Gln	He	He	His
			340					345					350		
Thr	Gly	Glu	Lys	Pro	Tyr	Gln	Cys	Asp	Ile	Cys	Gly	Lys	Val	Phe	Arg
		355					360					365			
Gln	Asn	Ser	Asn	Leu	Val	Asn	His	Gln	Arg	He	His	Thr	Gly	Glu	Lys
	370					375					380				
Pro	Tyr	Lys	Cys	Asn	He	Cys	Gly	Lys	Ser	Phe	Ser	G1n	Ser	Ser	
385					390					395					400
Leu	Ala	Thr	His		Thr	Val	His	Ser		Asn	Lys	Pro	Tyr	Lys	Cys
				405					410		_			415	
Asp	Glu	Cys		Lys	Thr	Phe	Lys		Ser	Ser	Ser	Leu		Thr	His
0.3	** 3	7.1	420	an.	0.1	0.1	,	425 D	Tr.	TT1	0		430	0	
GIn	Val		His	Thr	Gly	Glu		Pro	lyr	Ihr	Cys		vai	Cys	Asp
	17 1	435	C	C1	Α.	C .	440	1	۸1.	A	11.2	445	A	C1	11: -
Lys		Pne	ser	Gin	Arg			Leu	Ala	Arg		UIN	Arg	Gly	mis
Tl	450	C1	Lua	Dwo	T	455		Aan	C1	Cua	460	Lva	Vol	Dho	Sor
	GIY	61u	Lys	Pro		Lys	Cys	ASII	610	475	GIŸ	Lys	vai	Phe	480
465	The	San	Ша	Lov	470 Val	Cl.	Hic	Ara	Arc		Hic	The	61.	Glu	
OIU	HIL	ser	1115	485	val	GIY	111.5	ия	490		111.5	1111	019	Glu 495	Lys
Pro	Tvr	lve	Cvs		Lve	Cvs	Glv	l.vs			Lvs	Gln	Glv	Ser	Leu
110	4 V.I	L 7 3	U Y .3	1100		~ 1 .	- U 1 V	ت ربی		4 110			~ × ×		~~~~

			500					505					510		
Leu	Thr	Arg	His	Lys	lle	11e	His	Thr	Arg	Glu	Lys	Arg	Tyr	Gln	Cys
		515					520					525			
Gly	Glu	Cys	Gly	Lys	Val	Phe	Ser	Glu	Asn	Ser	Cys	Leu	Val	Arg	His
	530					535					540				
Leu	Arg	lle	llis	Thr	Gly	Glu	G1n	Pro	Tyr	Lys	Cys	Asn	Val	Cys	Gly
545					550					555					560
Lys	Val	Phe	Asn	Tyr	Ser	Gly	Asn	Leu	Ser	Ile	His	Lys	Arg	lle	Arg
				565					570					575	
Thr	Gly	Glu	Lys	Pro	Phe	Gln	Cys	Asn	Glu	Cys	Gly	Thr	Val	Phe	Arg
			580					585					590		
Asn	Tyr	Ser	Cys	Leu	Ala	Arg	His	Leu	Arg	He	llis	Thr	Gly	G1n	Lys
		595					600					605			
Pro	Tyr	Lys	Cys	Asn	Val	Cys	Gly	Lys	Val	Phe	Asn	Asp	Ser	G1 y	Asn
	610					615					620				
Leu	Ser	Asn	His	Lys	Arg	He	His	Thr	Gly	Glu	Lys	Pro	Phe	Gln	Cys
625					630					635					640
Asn	Glu	Cys	Gly	Lys	Va]	Phe	Ser	Tyr	Tyr	Ser	Cys	Leu	Ala	Arg	His
				645					650					655	
Arg	Lys	lle	llis	Ala	Gly	Glu	Lys	Pro	Tyr	Lys	Cys	Asn	Asp	Cys	Gly
			660					665					670		
Lys	Ala	Tyr	Thr	Gln	Arg	Ser	Ser	Leu	Thr	Lys	His	Leu	Ile	He	His
		675					680					685			
Thr	Gly	Glu	Lys	Pro	Tyr	Asn	Cys	Asn	Glu	Phe	Gly	Gly	Ala	Phe	He
	690					695					700				
Gln	Ser	Ser	Lys	Leu	Ala	Arg	Tyr	His	Arg	Asn	Pro	Thr	Gly	Glu	Lys
705					710					715					720
Pro	His	Lys	Cys	Ser	His	Cys	Gly	Arg	Thr	Phe	Ser	His	lle	Thr	Gly
				725					730					735	
Leu	Thr	Tyr	His	Gln	Arg	Arg	His	Thr	G1 y	Glu	Met	Pro	Tyr	Lys	Cys
			740					745					750		
He	Glu	Cys	Gly	Gln	Val	Phe	Asn	Ser	Thr	Ser	Asn	Leu	Ala	Arg	His
		755					760					765			
Arg	Arg	Ile	His	Thr	Gly	Glu	Lys	Pro	Tyr	Lys	Cys	Asn	Glu	Cys	Gly
	770					775					780				
Lys	Val	Phe	Arg	His	Gln	Ser	Thr	Leu	Ala	Arg	His	Arg	Ser	He	His

800 785 790 795 Thr Gly Glu Lys Pro Tyr Val Cys Ser Glu Cys Gly Lys Ala Phe Arg 805 810 Val Arg Ser Ile Leu Val Asn His Gln Lys Met His Thr Gly Asp Lys 820 830 825 Pro Tyr Lys Cys Asn Glu Cys Gly Lys Ala Phe Ile Glu Arg Ser Lys 840 Leu Val Tyr His Gln Arg Asn His Thr Gly Glu Lys Pro Tyr Lys Cys 850 855 860 Ile Glu Cys Gly Lys Ala Phe Gly Arg Phe Ser Cys Leu Asn Lys His 870 875 880 Gln Met Ile His Ser Gly Glu Lys Pro Tyr Lys 885

<210> 4836

<211> 374

<212> PRT

<213> Homo sapiens

<400> 459

Met Ile Glu Tyr Gln Ile Pro Val Ser Phe Lys Asp Val Val Gly
1 5 10 15

Phe Thr Gln Glu Glu Trp His Arg Leu Ser Pro Ala Gln Arg Ala Leu 20 25 30

Tyr Arg Asp Val Met Leu Glu Thr Tyr Ser Asn Leu Ala Ser Val Gly
35 40 45

Tyr Glu Gly Thr Lys Pro Asp Val Ile Leu Arg Leu Glu Glu Glu 50 55 60

Ala Pro Trp lle Gly Glu Ala Ala Cys Pro Gly Cys His Cys Trp Glu
65 70 75 80

Asp lle Trp Arg Val Asn lle Gln Arg Lys Arg Arg Gln Asp Met Leu 85 .90 95

Leu Arg Pro Gly Ala Ala lle Ser Lys Lys Thr Leu Pro Lys Glu Lys 100 105 110

Ser Cys Glu Tyr Asn Lys Phe Gly Lys Ile Ser Leu Leu Ser Thr Asp

		115					120					125			
Leu	Phe	Ser	Ser	lle	Gln	Ser	Pro	Ser	Asn	Trp	Asn	Pro	Cys	Gly	Lys
	130					135					140				
Asn	Leu	Asn	His	Asn	Leu	Asp	Leu	He	Gly	Phe	Lys	Arg	Asn	Cys	Ala
145					150					155					160
Lys	Lys	Gln	Asp	Glu	Cys	Tyr	Ala	Tyr	Gly	Lys	Leu	Leu	Gln	Arg	Ile
				165					170					175	
Asn	His	Gly	Arg	Arg	Pro	Asn	Gly	Glu	Lys	Pro	Arg	Gly	Cys	Ser	His
			180					185					190		
Cys	Glu	Lys	Ala	Phe	Thr	Gln	Asn	Pro	Ala	Leu	Met	Tyr	Lys	Pro	Ala
		195					200					205			
Val	Ser	Asp	Ser	Leu	Leu	Tyr	Lys	Arg	Lys	Arg	Val	Pro	Pro	Thr	Glu
	210					215					220				
Lys	Pro	His	Val	Cys	Ser	Glu	Cys	Gly	Lys	Ala	Phe	Cys	Tyr	Lys	Ser
225					230					235					240
Glu	Phe	lle	Arg	His	Gln	Arg	Ser	His	Thr	Gly	Glu	Lys	Pro	Tyr	Gly
				245					250					255	
Cys	Thr	Asp	Cys	Gly	Lys	Ala	Phe	Ser	His	Lys	Ser	Thr	Leu	Ile	Lys
			260					265					270		
His	Gln	Arg	lle	His	Thr	Gly	Val	Arg	Pro	Phe	Glu	Cys	Phe	Phe	Cys
		275					280					285			
Gly	Lys	Ala	Phe	Thr	Gln	Lys	Ser	His	Arg	Thr	Glu	His	Gln	Arg	Thr
	290					295					300				
His	Thr	Gly	Glu	Arg	Pro	Phe	Val	Cys	Ser	Glu	Cys	Gly	Lys	Ser	Phe
305					310					315					320
G1 y	Glu	Lys	Ser	Tyr	Leu	Asn	Val	His	Arg	Lys	Met	His	Thr	Gly	Glu
				325					330					335	
Arg	Pro	Tyr	Arg	Cys	Arg	Glu	Cys	G1y	Lys	Ser	Phe	Ser	Gln	Lys	Ser
			340					345					350		
Cys	Leu	Asn	Lys	His	Trp	Λrg	Thr	His	Phe	Gly	Glu	Ser	Ser	Leu	Arg
		355					360					365			
Ser	Lys	Ser	Ser	Asn	Thr										
	370														

```
<211> 896
<212> PRT
<213> Homo sapiens
<400> 460
Met Thr Glu Glu Ser Glu Glu Thr Val Leu Tyr Ile Glu His Arg Tyr
                                     10
Val Cys Ser Glu Cys Asn Gln Leu Tyr Gly Ser Leu Glu Glu Val Leu
             20
                                 25
                                                      30
Met His Gln Asn Ser His Val Pro Gln Gln His Phe Glu Leu Val Gly
         35
                             40
Val Ala Asp Pro Gly Val Thr Val Ala Thr Asp Thr Ala Ser Gly Thr
                         55
Gly Leu Tyr Gln Thr Leu Val Gln Glu Ser Gln Tyr Gln Cys Leu Glu
 65
                     70
                                         75
Cys Gly Gln Leu Leu Met Ser Pro Ser Gln Leu Leu Glu His Gln Glu
                                     90
Leu His Leu Lys Met Met Ala Pro Gln Glu Ala Val Pro Ala Glu Pro
                                                     110
            100
                                105
Ser Pro Lys Ala Pro Pro Leu Ser Ser Ser Thr 11e His Tyr Glu Cys
                            120
        115
Val Asp Cys Lys Ala Leu Phe Ala Ser Gln Glu Leu Trp Leu Asn His
                        135
                                             140
Arg Gln Thr His Leu Arg Ala Thr Pro Thr Lys Ala Pro Ala Pro Val
145
                    150
                                         155
                                                             160
Val Leu Gly Ser Pro Val Val Leu Gly Pro Pro Val Gly Gln Ala Arg
                165
                                    170
Val Ala Val Glu His Ser Tyr Arg Lys Ala Glu Glu Gly Glu Gly
                                185
                                                     190
            180
Ala Thr Val Pro Ser Ala Ala Ala Thr Thr Glu Val Val Thr Glu
                                                 205
        195
                            200
Val Glu Leu Leu Tyr Lys Cys Ser Glu Cys Ser Gln Leu Phe Gln
                        215
Leu Pro Ala Asp Phe Leu Glu His Gln Ala Thr His Phe Pro Ala Pro
225
                    230
                                         235
                                                             240
Val Pro Glu Ser Gln Glu Pro Ala Leu Gln Gln Glu Val Gln Ala Ser
```

				245					250					255	
Ser	Pro	Ala	Glu	Val	Pro	Val	Ser	Gln	Pro	Asp	Pro	Leu	Pro	Ala	Ser
			260					265					270		
Asp	His	Ser	Tyr	Glu	Leu	Arg	Asn	Gly	Glu	Ala	Ile	Gly	Arg	Asp	Arg
		275					280					285			
Arg	Gly	Arg	Arg	Ala	Arg	Arg	Asn	Asn	Ser	Gly	Glu	Ala	Gly	Gly	Ala
	290					295					300				
Ala	Thr	Gln	Glu	Leu	Phe	Cys	Ser	Ala	Cys	Asp	Gln	Leu	Phe	Leu	Ser
305					310					315					320
Pro	His	Gln	Leu	Gln	Gln	His	Leu	Arg	Ser	His	Arg	Glu	G1 y	Val	Phe
				325					330					335	
Lys	Cys	Pro	Leu	Cys	Ser	Arg	Val	Phe	Pro	Ser	Pro	Ser	Ser	Leu	Asp
			340					345					350		
Gln	His	Leu	Gly	Asp	His	Ser	Ser	Glu	Ser	His	Phe	Leu	Cys	Val	Asp
		355					360					365			
Cys	Gly	Leu	Ala	Phe	Gly	Thr	Glu	Ala	Leu	Leu	Leu	Ala	His	Arg	Arg
	370					375					380				
Ala	His	Thr	Pro	Asn	Pro	Leu	His	Ser	Cys	Pro	Cys	Gly	Lys	Thr	Phe
385					390					395					400
Val	Asn	Leu	Thr	Lys	Phe	Leu	Tyr	His	Arg	Arg	Thr	His	Gly	Val	Gly
				405					410					415	
Gly	Val	Pro	Leu	Pro	Thr	Thr	Pro	Val	Pro	Pro	Glu	Glu	Pro	Val	lle
			420					425					430		
Gly	Phe	Pro	Glu	Pro	Ala	Pro	Ala	Glu	Thr	Gly	Glu	Pro	Glu	Ala	Pro
		435					440					445			
Glu	Pro	Pro	Val	Ser	Glu	Glu	Thr	Ser	Ala	Gly	Pro	Ala	Ala	Pro	Gly
	450		_				_				460				0.1
	Tyr	Arg	Cys	Leu		Cys	Ser	Arg	Glu		Gly	Lys	Ala	Leu	
465				0.1	470	-				475	0.7				480
Leu	Thr	Arg	His		Arg	Phe	Val	HIS		Leu	Glu	Arg	Arg		Lys
0	C			485			121	,	490	1	C -	117 .	V . 1	495	۸
Cys	Ser	He	-	Gly	Lys	Met	Phe		Lys	Lys	Ser	HIS		Arg	Asn
Δ	1	Λ	500	11.2	T1	C1.	C1	505	Das	Dh	Dass	Cus	510 Pro	100	Cuc
Arg	ren		mr	птѕ	ınr	Gly		нгg	L1.0	r116	1.10	525	1.10	лsр	CyS
C ~ 1-	Luc	515 Pro	Dha	100	Ç 0.1-	Pro	520	Acr	Lou	Ala	A = a		Λκα	Lou	The
> O 1	1 17 0	11111	1.170	Hen	>0 1°	rra	1113	acn	L (C) 1 1	/A I (3)	44 T (T	1111	73 E CT	1 (2))	1.617

	530					535					540				
His	Thr	Gly	Glu	Arg	Pro	Tyr	Arg	Cys	Gly	Asp	Cys	Gly	Lys	Ala	Phe
545					550					555					560
Thr	Gln	Ser	Ser	Thr	Leu	Arg	Gln	His	Arg	Leu	Val	His	Ala	Gln	His
				565					570					575	
Phe	Pro	Tyr	Arg	Cys	Gln	Glu	Cys	Gly	Val	Arg	Phe	His	Arg	Pro	Tyr
			580					585					590		
Arg	Leu	Leu	Met	His	Arg	Tyr	His	His	Thr	Gly	Glu	Tyr	Pro	Tyr	Lys
		595					600					605			
Cys	Arg	Glu	Cys	Pro	Arg	Ser	Phe	Leu	Leu	Arg	Arg	Leu	Leu	Glu	Val
	610					615					620				
His	Gln	Leu	Val	Val	His	Ala	Gly	Arg	Gln	Pro	His	Arg	Cys	Pro	Ser
625					630					635					640
Cys	Gly	Ala	Ala	Phe	Pro	Ser	Ser	Leu	Arg	Leu	Arg	Glu	His	Arg	Cys
				645					650					655	
Ala	Ala	Ala	Ala	Ala	Gln	Ala	Pro	Arg	Arg	Phe	Glu	Cys	Gly	Thr	Cys
			660					665					670		
Gly	Lys	Lys	Val	Gly	Ser	Ala	Ala	Arg	Leu	·G1n	Ala	His	Glu	Ala	Ala
		675					680					685			
His	Ala	Ala	Ala	Gly	Pro	Gly	Glu	Val	Leu	Ala	Lys	Glu	Pro	Pro	Ala
	690					695					700				
Pro	Arg	Ala	Pro	Arg	Ala	Thr	Arg	Ala	Pro	Val	Ala	Ser	Pro	Ala	Ala
705					710					715					720
Leu	Gly	Ser	Thr	Ala	Thr	Ala	Ser	Pro	Ala	Ala	Pro	Ala	Arg	Arg	Arg
				725					730					735	
Gly	Leu	G1u	Cys	Ser	Glu	Cys	Lys	Lys	Leu	Phe	Ser	Thr	Glu	Thr	Ser
			740					745					750		
Leu	Gln	Val	His	Arg	Arg	He	His	Thr	Gly	Glu	Arg	Pro	Tyr	Pro	Cys
		755					760					765			
Pro	Asp	Cys	Gly	Lys	Ala	Phe	Arg	Gln	Ser	Thr	His	Leu	Lys	Asp	His
	770					775					780				
Arg	Arg	Leu	His	Thr		G]u	Arg	Pro	Phe		Cys	Glu	Val	Cys	
785					790					795					800
Lys	Ala	Phe	Ala	He	Ser	Met	Arg	Leu		Glu	His	Arg	Arg		His
				805					810					815	

Thr Gly Glu Arg Pro Tyr Ser Cys Pro Asp Cys Gly Lys Ser Tyr Arg Ser Phe Ser Asn Leu Trp Lys His Arg Lys Thr His Gln Gln His Gln Ala Ala Val Arg Gln Gln Leu Ala Glu Ala Glu Ala Val Gly Leu Ala Val Met Glu Thr Ala Val Glu Ala Leu Pro Leu Val Glu Ala lle Glu Ile Tyr Pro Leu Ala Glu Ala Glu Gly Val Gln Ile Ser Gly

<210> 4838

<211> 464

<212> PRT

<213> Homo sapiens

<400> 461

Met Phe Glu Asn Glu Ser Arg Lys Ile Phe Ser Glu Met Pro Glu Gly Glu Ser Ala Gln His Ser Asp Gly Glu Ser Asp Phe Glu Arg Asp Ala Gly lle Gln Arg Leu Gln Gly His Thr Pro Gly Glu Asp His Gly Glu Val Val Ser Gln Asp Arg Glu Val Gly Gln Leu Ile Gly Leu Gln Gly Thr Tyr Leu Gly Glu Lys Pro Tyr Glu Cys Pro Gln Cys Gly Lys Thr Phe Ser Pro Lys Ser His Leu Ile Thr His Glu Arg Thr His Thr Gly Glu Lys Tyr Tyr Lys Cys Asp Glu Cys Gly Lys Ser Phe Ser Asp Gly Ser Asn Phe Ser Arg His Gln Thr Thr His Thr Gly Glu Lys Pro Tyr Lys Cys Arg Asp Cys Gly Lys Ser Phe Ser Arg Ser Ala Asn Leu lle

	His	Gln	Arg	He	His	Thr	Gly	Glu	Lys		Phe	Gln	Cys	Ala	Glu
145					150					155					160
Cys	G1 y	Lys	Ser	Phe	Ser	Arg	Ser	Pro	Asn	Leu	He	Λla	His	Gln	Arg
				165					170					175	
Thr	His	Thr	Gly	Glu	Lys	Pro	Tyr	Ser	Cys	Pro	Glu	Cys	Gly	Lys	Ser
			180					185					190		
Phe	Gly	Asn	Arg	Ser	Ser	Leu	Asn	Thr	His	G1n	Gly	He	His	Thr	Gly
		195					200					205			
Glu	Lys	Pro	Tyr	Glu	Cys	Lys	Glu	Cys	Gly	Glu	Ser	Phe	Ser	Tyr	Asn
	210					215					220				
Ser	Asn	Leu	He	Arg	His	Gln	Arg	He	His	Thr	G1 y	Glu	Glu	Pro	Tyr
225					230					235					240
Lys	Cys	Thr	Asp	Cys	Gly	Gln	Arg	Phe	Ser	Gln	Ser	Ser	Ala	Leu	lle
				245					250					255	
Thr	His	Arg	Arg	Thr	His	Thr	Gly	Glu	Lys	Pro	Tyr	Gln	Cys	Ser	Glu
			260					265					270		
Cys	Gly	Lys	Ser	Phe	Ser	Arg	Ser	Ser	Asn	Leu	Ala	Thr	His	Arg	Arg
		275					280					285			
Thr	His	Met	Val	Glu	Lys	Pro	Tyr	Lys	Cys	Gly	Val	Cys	Gly	Lys	Ser
	290					205					200				
	250					295					300				
Phe		Gln	Ser	Ser	Ser		lle	Ala	His	Gln		Met	His	Thr	Gly
Phe 305		Gln	Ser	Ser	Ser 310		lle	Ala	His	Gln 315		Met	His	Thr	Gly 320
305	Ser				310	Leu				315	G1 y	Met Phe			320
305	Ser				310	Leu				315	G1 y				320
305 Glu	Ser Lys	Pro	Tyr	G1u 325	310 Cys	Leu Leu	Thr	Cys	Gly 330	315 Glu	Gly Ser		Ser	Trp 335	320 Ser
305 Glu	Ser Lys	Pro	Tyr	G1u 325	310 Cys	Leu Leu	Thr	Cys	Gly 330	315 Glu	Gly Ser	Phe	Ser	Trp 335	320 Ser
305 Glu Ser	Ser Lys Asn	Pro Leu	Tyr Leu 340	Glu 325 Lys	310 Cys His	Leu Leu Gln	Thr Arg	Cys Ile 345	Gly 330 His	315 Glu Thr	Gly Ser Gly	Phe	Ser Lys 350	Trp 335 Pro	320 Ser Tyr
305 Glu Ser	Ser Lys Asn	Pro Leu	Tyr Leu 340	Glu 325 Lys	310 Cys His	Leu Leu Gln	Thr Arg	Cys Ile 345	Gly 330 His	315 Glu Thr	Gly Ser Gly	Phe Glu	Ser Lys 350	Trp 335 Pro	320 Ser Tyr
305 Glu Ser Lys	Ser Lys Asn Cys	Pro Leu Ser 355	Tyr Leu 340 Glu	Glu 325 Lys Cys	310 Cys His	Leu Leu Gln Lys	Thr Arg Cys 360	Cys Ile 345 Phe	Gly 330 His Ser	315 Glu Thr Gln	Gly Ser Gly Arg	Phe Glu Ser	Ser Lys 350 GIn	Trp 335 Pro Leu	320 Ser Tyr Val
305 Glu Ser Lys	Ser Lys Asn Cys	Pro Leu Ser 355	Tyr Leu 340 Glu	Glu 325 Lys Cys	310 Cys His	Leu Leu Gln Lys	Thr Arg Cys 360	Cys Ile 345 Phe	Gly 330 His Ser	315 Glu Thr Gln	Gly Ser Gly Arg	Phe Glu Ser 365	Ser Lys 350 GIn	Trp 335 Pro Leu	320 Ser Tyr Val
305 Glu Ser Lys Val	Ser Lys Asn Cys His 370	Pro Leu Ser 355 GIn	Tyr Leu 340 Glu Arg	Glu 325 Lys Cys	310 Cys His Gly	Leu Gln Lys Thr 375	Thr Arg Cys 360 Gly	Cys Ile 345 Phe Glu	Gly 330 His Ser Lys	315 Glu Thr Gln Pro	Gly Ser Gly Arg Tyr 380	Phe Glu Ser 365	Ser Lys 350 Gln Cys	Trp 335 Pro Leu Leu	320 Ser Tyr Val Met
305 Glu Ser Lys Val	Ser Lys Asn Cys His 370	Pro Leu Ser 355 GIn	Tyr Leu 340 Glu Arg	Glu 325 Lys Cys	310 Cys His Gly	Leu Gln Lys Thr 375	Thr Arg Cys 360 Gly	Cys Ile 345 Phe Glu	Gly 330 His Ser Lys	315 Glu Thr Gln Pro	Gly Ser Gly Arg Tyr 380	Phe Glu Ser 365 Lys	Ser Lys 350 Gln Cys	Trp 335 Pro Leu Leu	320 Ser Tyr Val Met
305 Glu Ser Lys Val Cys 385	Lys Asn Cys His 370 Gly	Pro Leu Ser 355 GIn Lys	Tyr Leu 340 Glu Arg Ser	Glu 325 Lys Cys Thr	310 Cys His Gly His Ser 390	Leu Gln Lys Thr 375 Arg	Thr Arg Cys 360 Gly Gly	Cys Ile 345 Phe Glu Ser	Gly 330 His Ser Lys	315 Glu Thr Gln Pro Leu 395	Gly Ser Gly Arg Tyr 380 Val	Phe Glu Ser 365 Lys	Ser Lys 350 GIn Cys	Trp 335 Pro Leu Leu	320 Ser Tyr Val Met Arg 400
305 Glu Ser Lys Val Cys 385	Lys Asn Cys His 370 Gly	Pro Leu Ser 355 GIn Lys	Tyr Leu 340 Glu Arg Ser	Glu 325 Lys Cys Thr	310 Cys His Gly His Ser 390	Leu Gln Lys Thr 375 Arg	Thr Arg Cys 360 Gly Gly	Cys Ile 345 Phe Glu Ser	Gly 330 His Ser Lys	315 Glu Thr Gln Pro Leu 395	Gly Ser Gly Arg Tyr 380 Val	Phe Glu Ser 365 Lys Met	Ser Lys 350 GIn Cys	Trp 335 Pro Leu Leu	320 Ser Tyr Val Met Arg 400
305 Glu Ser Lys Val Cys 385 Ala	Lys Asn Cys His 370 Gly	Pro Leu Ser 355 GIn Lys	Tyr Leu 340 Glu Arg Ser Gly	G1u 325 Lys Cys Thr Phe Asp 405	310 Cys His Gly His Ser 390 Lys	Leu Gln Lys Thr 375 Arg	Thr Arg Cys 360 Gly Gly Tyr	Cys Ile 345 Phe Glu Ser	Gly 330 His Ser Lys He Cys 410	315 Glu Thr Gln Pro Leu 395 Pro	Gly Ser Gly Arg Tyr 380 Val	Phe Glu Ser 365 Lys Met	Ser Lys 350 GIn Cys His	Trp 335 Pro Leu Leu Gln Lys 415	320 Ser Tyr Val Met Arg 400 Gly

Glu Lys Pro Tyr Lys Cys Pro Glu Cys Gly Lys Gly Phe Ser Asn Ser
435

440

445

Ser Asn Phe Ile Thr His Gln Arg Thr His Met Lys Glu Lys Leu Tyr
450

455

460

<210> 4839

<211> 347

<212> PRT

<213> Homo sapiens

<400> 462

Met Ser Gly Pro Gly Asn Lys Arg Ala Ala Gly Asp Gly Gly Ser Gly
1 5 10 15

Pro Pro Glu Lys Lys Leu Ser Arg Glu Glu Lys Thr Thr Thr Leu
20 25 30

Ile Glu Pro 1le Arg Leu Gly Gly 1le Ser Ser Thr Glu Glu Met Asp
35 40 45

Leu Lys Val Leu Gln Phe Lys Asn Lys Lys Leu Ala Glu Arg Leu Glu 50 55 60

Gln Arg Gln Ala Cys Glu Asp Glu Leu Arg Glu Arg Ile Glu Lys Leu 65 70 75 80

Glu Lys Arg Gln Ala Thr Asp Asp Ala Thr Leu Leu lle Val Asn Arg

85 90 95

Tyr Trp Ala Gln Leu Asp Glu Thr Val Glu Ala Leu Leu Arg Cys His 100 105 110

Glu Ser Arg Ile Arg Glu Leu Glu Glu Arg Asp Arg Glu Ser Lys
115 120 125

Lys lle Ala Asp Glu Asp Ala Leu Arg Arg lle Arg Gln Ala Glu Glu 130 135 140

Gln Ile Glu His Leu Gln Arg Lys Leu Gly Ala Thr Lys Gln Glu Glu 145 150 155 160

Glu Ala Leu Leu Ser Glu Met Asp Ala Gln Leu Leu Thr Val Gln Lys 165 170 175

Leu Glu Glu Lys Glu Arg Ala Leu Gln Gly Ser Leu Gly Gly Val Glu 180 185 190

Lys	Glu	Leu	Thr	Leu	Arg	Ser	Gln	Ala	Leu	Glu	Leu	Asn	Lys	Arg	Lys
		195					200					205			
Ala	Val	Glu	Ala	Ala	Gln	Leu	Ala	Glu	Asp	Leu	Lys	Val	Gln	Leu	Glu
	210					215					220				
His	Val	Gln	Thr	Arg	Leu	Arg	Glu	He	Gln	Pro	Cys	Leu	Ala	Glu	Ser
225					230					235					240
Arg	Λla	Ala	Arg	Glu	Lys	Glu	Ser	Phe	Asn	Leu	Lys	Arg	Ala	Gln	Glu
				245					250					255	
Asp	lle	Ser	Arg	Leu	Arg	Arg	Lys	Leu	Glu	Lys	Gln	Arg	Lys	Val	Glu
			260					265					270		
Val	Tyr	Ala	Asp	Ala	Asp	Glu	lle	Leu	Gln	Glu	Glu	lle	Lys	Glu	Tyr
		275					280					285			
Lys	Ala	Arg	Leu	Thr	Cys	Pro	Cys	Cys	Asn	Thr	Arg	Lys	Lys	Asp	Ala
	290					295					300				
Val	Leu	Thr	Lys	Cys	Phe	His	Val	Phe	Cys	Phe	Glu	Cys	Val	Arg	Gly
305					310					315					320
Arg	Tyr	Glu	Ala	Arg	Gln	Arg	Lys	Cys	Pro	Lys	Cys	Asn	Ala	Ala	Phe
				325					330					335	
Gly	Ala	His	Asp	Phe	His	Arg	Ile	Tyr	He	Ser					
			340					345							

<210> 4840

<211> 997

<212> PRT

<213> Homo sapiens

<400> 463

 Met
 Glu
 Ala
 Ser
 Trp
 Arg
 Gln
 Val
 Ala
 Gly
 Gly
 Arg
 Gly
 Arg
 Ser
 Arg

 1
 5
 5
 10
 10
 15
 15

 Gly
 Arg
 Ala
 Ala
 Pro
 Ser
 Gly
 Asn
 Gly
 Val
 His
 Leu
 Arg
 Gly

 Ala
 Gly
 Gly
 Arg
 Glu
 Lys
 Gly
 Ser
 Val
 Gly
 Ala
 Val
 Pro
 Ser
 Gly

 Thr
 Ser
 Pro
 Gly
 Gly
 Val
 Ala
 Thr
 Ala
 Ala
 Ala
 Ala
 Gly
 Ser
 Arg
 His

 50
 55
 55
 60
 Fro
Ser 65	Pro	Ala	Gly	Ser	G1n 70	Ala	Leu	Gln	Thr	Thr 75	Ala	Ala	Ser	Glu	Leu 80
		C1	1	1		C1	C1	11.	1		A T	Aan	C1	A 1 a	
Met	ser	GIR	Lys	Lys 85	rne	GIU	GIU	116	90	Lys	Ма	ASII	om	95	MIa
Ala	Arg	Lys	Leu	Val	Glu	Glu	G1n	Phe	Ser	Ser	Ser	Ser	Glu	Glu	Gly
			100					105					110		
Asp	Glu	Asp	Phe	Glu	G1 v	Lvs	Gln	Glv	Lvs	He	Leu	Ala	Asn	Thr	Phe
•		115			•	•	120		·			125			
Ile	Thr		Thr	Thr	Gln	Thr	Asp	Gly	Asp	Thr	Arg	Glu	Leu	Glu	Arg
	130	•				135	•				140				
Thr		Gln	Tvr	Val	Asn		Ala	Phe	Gln	Ala		Ala	Met	Thr	Cys
145			- 3 -		150					155	٠				160
	11e	Cvs	He	Ala		Val	Lvs	Arg	Asn		Ala	Val	Trp	Ser	
		- ,		165			3	. 0	170				•	175	•
Ser	Glv	Cvs	Phe	Cys	Ile	Phe	His	Met	Pro	Cvs	lle	Gln	Lvs	Trp	Ala
		- 3	180	•				185		•			190	•	
Lvs	Asp	Ser		Phe	Leu	Val	Ser		Val	Thr	Asp	Asp	Asp	Phe	Gly
-,-		195					200				•	205	•		•
Lvs	Lvs		Cvs	Pro	Trp	Pro		Pro	Lys	Cvs	Arg		Glu	Tyr	Lys
-,-	210		-,-			215	-,-			- , -	220			,	•
Arg		Glu	Thr	Pro	Ser		Tvr	Tvr	Cvs	Tvr		Gly	Lvs	Val	Glu
Ü						Ŭ	•	-	•	·	•	-	•		
225					230					235					240
Asp	Pro	Pro	Leu	Asp	Pro	Trp	Leu	Val	Pro	His	Ser	Cys	Gly	Gln	Val
				245					250					255	
Cys	Glu	Arg	Glu	Phe	Lys	Pro	Pro	Cys	Gly	His	Lys	Cys	Leu	Leu	Leu
			260					265					270		
Cys	His	Pro	Gly	Pro	Cys	Pro	Pro	Cys	Pro	Lys	Met	Val	Thr	Thr	Thr
		275					280					285			
Cys	Tyr	Cys	Lys	Lys	Ala	Lys	Pro	He	Pro	Arg	Λrg	Cys	Ser	Ala	Lys
	290					295					300				
Glu	Trp	Ser	Cys	G1n	Leu	Pro	Cys	Gly	G1n	Lys	Leu	Leu	Cys	Gly	Gln
305					310					315					320
	Lys	Cys	Glu	Asn	Pro	Cys	His	Ala	Gly	Ser	Cys	G]n	Pro	Cys	Pro
				325					330					335	
Arg	Val	Ser	Arg	Gln	Lys	Cys	Val	Cys	Gly	Lys	Lys	Val	Ala	Glu	Arg

			340					345					350		
Ser	Cys	Ala	Ser	Pro	Leu	Trp	His	Cys	Asp	G1n	Val	Cys	Gly	Lys	Thr
		355					360					365			
Leu	Pro	Cys	Gly	Asn	His	Thr	Cys	Glu	Gln	Val	Cys	His	Val	Gly	Ala
	370	•				375					380				
Cys	Gly	Glu	Cys	Pro	Arg	Ser	Gly	Lys	Arg	Phe	Cys	Pro	Cys	Gln	Lys
385					390					395					400
Ser	Lys	Phe	Ser	Leu	Pro	Cys	Thr	Glu	Asp	Val	Pro	Thr	Cys	Gly	Asp
				405					410					415	
Ser	Cys	Asp	Lys	Val	Leu	Glu	Cys	Gly	Ile	His	Arg	Cys	Ser	Gln	Arg
			420					425					430		
Cys	His	Arg	Gly	Pro	Cys	Glu	Thr	Cys	Arg	Gln	Glu	Val	Glu	Lys	His
		435					440					445			
Cys	Arg	Cys	Gly	Lys	His	Thr	Lys	Arg	Met	Pro	Cys	His	Lys	Pro	Tyr
	450					455					460				
Leu	Cys	Glu	Thr	Lys	Cys	Val	Lys	Met	Arg	Asp	Cys	Gln	Lys	His	Gln
465					470					475					480
Cys	Arg	Arg	Lys	Cys	Cys	Pro	Gly	Asn	Cys	Pro	Pro	Cys	Asp	Gln	Asn
				485					490					495	
Cys	Gly	Arg	Thr	Leu	Gly	Cys	Arg	Asn	His	Lys	Cys	Pro	Ser	Val	Cys
			500					505					510		
His	Arg	Gly	Ser	Cys	Tyr	Pro	Cys	Pro	Glu	Thr	Val	Asp	Val	Lys	Cys
		515					520					525			
Asn		Gly	Asn	Thr	Lys	Va]	Thr	Va]	Pro	Cys	G] y	Arg	Glu	Arg	Thr
	530					535					540				
	Arg	Pro	Pro	Lys	Cys	Lys	Glu	Gln	Cys	Ser	Arg	Pro	Pro	Thr	
545					550					555					560
His	His	Thr	Ser		Glu	Lys	His	Arg		His	Phe	Gly	Ser		Pro
				565					570					575	_
Pro	Cys	His		Pro	Cys	Gln	Lys		Leu	Glu	Lys	Cys		His	Leu
			580				0.7	585					590		
Cys	***		15	_										0.1	
	Pro		Pro	Cys	His	Asp		Ala	Leu	He	Lys		Thr	Gly	Arg
11.		595					600					605			
His	Gln	595				Trp	600				Glu	605			
	Gln 610	595 Pro	Thr	Gly	Pro		600 Gl _u	Gln	Pro	Ser	Glu 620	605 Pro	Ala	Phe	He

625					630					635					640
Cys	Leu	Gly	Lys	His	$Gl\mathbf{u}$	Val	Ser	Pro	Leu	Pro	Cys	His	Ala	Val	Gly
				645					650					655	
Pro	Tyr	Ser	Cys	Lys	Arg	Val	Cys	Gly	Arg	lle	Leu	Asp	Cys	Gln	Asn
			660					665					670		
His	Thr	Cys	Met	Lys	Glu	Cys	His	Lys	Val	Thr	Lys	Thr	Asp	Gly	Cys
		675					680					685			
Thr	Gly	Lys	Asn	Lys	Ala	Gly	Pro	Glu	Cys	Leu	His	Cys	Glu	Glu	Gly
	690					695					700				
Cys	Ser	Lys	Ser	Arg	Pro	Leu	Gly	Cys	Leu	His	Pro	Cys	He	Leu	Arg
705					710					715					720
Cys	His	Pro	Gly	Glu	Cys	Pro	Pro	Cys	Val	Gln	Met	Leu	Arg	He	Lys
				725					730					735	
Cys	His	Cys	Lys]]e	Thr	Ser	Leu	Tyr	Val	Glu	Cys	Arg	Lys	He	Thr
			740					745					750		
Thr	Ala	Asp	Val	Asn	Glu	Lys	Asn	Leu	Leu	Ser	Cys	Cys	Lys	Asn	Gln
		755					760					765			
Cys	Pro	Lys	Glu	Leu	Pro	Cys	Gly	His	Arg	Cys	Lys	Glu	Met	Cys	His
	770					775					780				
	Gly	Glu	Cys	Pro		Asn	Cys	Asn	Gln		Val	Lys	Leu	Arg	
785					790					795					800
Pro	Cys	Lys	Arg		Lys	Lys	Glu	Leu		Cys	Asn	Lys	Val		Glu
				805					810					815	
Asn	Gln	Val		lle	Glu	Cys	Asp		Thr	Cys	Lys	Glu		Lys	Arg
_		_	820		_			825					830	0.1	0.1
Lys	Ala		Glu	He	Lys	Glu			Ala	Lys	Ala			Glu	Glu
0.1	,	835		61	0.1		840		C1		DI	845		. U	
Glu		Arg	Arg	61n	GIn	Ala	Glu	Leu	Glu	Ala		61u	Asn	Arg	Leu
1	850	A	11: -		C1	855	TL	C1	т	1	860	T!	V = 1	Tl	V = 1
	Inr	Asn	HIS	11e		His	Inr	61 y	iyr		Asn	ınr	vai	inr	
865	Dana	A	C1	Can	870	Cua	A16	Co.	C1	875	Luc	Aan	C1	Cln	880
ser	rro	ASP	GIY		rea	Cys	Ala	261		OTÀ	Lys	nsp	GIŸ		ита
Mo+	Lou	Trr	Acr	885	Acr	61	Glv	lve	890	Lou	Tun	The	Lou	895	61.
met	Leu	пþ	900	Leu	nSII	Glu	OTA	905	1112	ren	1 y 1	1.11.1	910	nsp	оту
Cl _v	Acr	Πο		Acn	Δ1α	Lou	Cvc		Sor	Pro	Acr	Ara		Trn	ىم ا

915			920			925		
Cys Ala Ala T	Thr Gly	Pro Ser	He Lys	Ile 1	Trp Asp	Leu G	lu Gly	Lys
930		935			940			
lle lle Val A	Asp Glu	Leu Lys	Gln Glu	Val 1	lle Ser	Thr S	er Ser	Lys
945		950		Ć	955			960
Ala Glu Pro F	Pro Gln	Cys Thr	Ser Leu	Ala	Trp Ser	Ala A		Gln
	965			970			975	
Thr Leu Phe A		Tyr Thr			Val Arg			Val
	980 Chan Ama		985			9	90	
Thr lle Gly T	inr Arg							
993								
			•					
<210> 4841								
<211> 709								
<212> PRT								
<213> Homo sa	apiens							
<400> 464								
Met Lys Arg (Ser Asp	Glu Leu		Gln Gln	Gln G		Glu
Met Lys Arg (5			10			15	
Met Lys Arg (5 Gly Leu		Ala Ala	10 Ser l			15 Gly Ala	
Met Lys Arg (l Asp Gly Ala (5 Gly Leu 20	Glu Asp	Ala Ala 25	10 Ser I	His Leu	Pro G	15 Gly Ala 30	Asp
Met Lys Arg C I Asp Gly Ala C Leu Arg Pro C	5 Gly Leu 20	Glu Asp	Ala Ala 25 Gly Ala	10 Ser I	His Leu	Pro G	15 Gly Ala 30	Asp
Met Lys Arg (1 Asp Gly Ala (Leu Arg Pro (35	5 Gly Leu 20 Gly Glu	Glu Asp	Ala Ala 25 Gly Ala 40	10 Ser I Asn S	His Leu Ser Ala	Pro G	15 Gly Ala 30 Gly Pro	Asp Thr
Met Lys Arg C I Asp Gly Ala C Leu Arg Pro C	5 Gly Leu 20 Gly Glu	Glu Asp	Ala Ala 25 Gly Ala 40	10 Ser I Asn S	His Leu Ser Ala	Pro G	15 Gly Ala 30 Gly Pro	Asp Thr
Met Lys Arg C I Asp Gly Ala C Leu Arg Pro C 35 Ser Asp Ala C	5 Gly Leu 20 Gly Glu Gly Ala	Glu Asp Thr Thr Ala Ala 55	Ala Ala 25 Gly Ala 40 Ala Pro	Ser I Asn S	His Leu Ser Ala Pro Gly 60	Pro G Gly G 45 Pro A	15 Gly Ala 30 Gly Pro	Asp Thr Lys
Met Lys Arg C I Asp Gly Ala C Leu Arg Pro C 35 Ser Asp Ala C	5 Gly Leu 20 Gly Glu Gly Ala	Glu Asp Thr Thr Ala Ala 55	Ala Ala 25 Gly Ala 40 Ala Pro	Ser I Asn S	His Leu Ser Ala Pro Gly 60	Pro G Gly G 45 Pro A	15 Gly Ala 30 Gly Pro	Asp Thr Lys
Met Lys Arg C I Asp Gly Ala C Leu Arg Pro C 35 Ser Asp Ala C 50 Pro Pro Asp I	5 Gly Leu 20 Gly Glu Gly Ala Leu Lys	Glu Asp Thr Thr Ala Ala 55 Lys Ile 70	Ala Ala 25 Gly Ala 40 Ala Pro	Ser I Asn S Asn I	His Leu Ser Ala Pro Gly 60 Ser Glu 75	Pro G Gly G 45 Pro A Gly S	15 Gly Ala 30 Gly Pro Arg Ser Ger Met	Asp Thr Lys Phe 80
Met Lys Arg C I Asp Gly Ala C Leu Arg Pro C 35 Ser Asp Ala C 50 Pro Pro Asp I 65 Gly His Gly I	5 Gly Leu 20 Gly Glu Gly Ala Leu Lys Eeu Lys 85	Glu Asp Thr Thr Ala Ala 55 Lys Ile 70 His Leu	Ala Ala 25 Gly Ala 40 Ala Pro Gln Glr	Ser I Asn S Asn I Leu S Ser I	His Leu Ser Ala Pro Gly 60 Ser Glu 75 Arg Arg	Pro G Gly G 45 Pro A Gly S Arg S	15 Gly Ala 30 Gly Pro Arg Ser Ger Met Ger Arg 95	Asp Thr Lys Phe 80 Glu
Met Lys Arg C I Asp Gly Ala C Leu Arg Pro C 35 Ser Asp Ala C 50 Pro Pro Asp I 65 Gly His Gly II Arg Glu His C	5 Gly Leu 20 Gly Glu Gly Ala Leu Lys Leu Lys 85 Gln Thr	Glu Asp Thr Thr Ala Ala 55 Lys Ile 70 His Leu	Ala Ala 25 Gly Ala 40 Ala Pro Gln Glr Phe His	Asn S Asn I Leu S Ser J Gln G	His Leu Ser Ala Pro Gly 60 Ser Glu 75 Arg Arg	Pro G Gly G 45 Pro A Gly S Gly S	15 Gly Ala 30 Gly Pro Arg Ser Ser Met Ser Arg 95 Gln Gln	Asp Thr Lys Phe 80 Glu
Met Lys Arg C I Asp Gly Ala C Leu Arg Pro C 35 Ser Asp Ala C 50 Pro Pro Asp I 65 Gly His Gly I Arg Glu His C	5 Gly Leu 20 Gly Glu Gly Ala Leu Lys 85 Gln Thr	Glu Asp Thr Thr Ala Ala 55 Lys Ile 70 His Leu Ser Gln	Ala Ala 25 Gly Ala 40 Ala Pro Gln Glr Phe His Asp Ser 105	Ser I Asn S Asn I Leu S Gln G	His Leu Ser Ala Pro Gly 60 Ser Glu 75 Arg Arg	Pro G Gly G 45 Pro A Gly S Gln G	I5 Gly Ala 30 Gly Pro Arg Ser Ger Met 95 Gln Gln	Asp Thr Lys Phe 80 Glu Gln
Met Lys Arg C I Asp Gly Ala C Leu Arg Pro C 35 Ser Asp Ala C 50 Pro Pro Asp I 65 Gly His Gly II Arg Glu His C	5 Gly Leu 20 Gly Glu Gly Ala Leu Lys 85 Gln Thr	Glu Asp Thr Thr Ala Ala 55 Lys Ile 70 His Leu Ser Gln	Ala Ala 25 Gly Ala 40 Ala Pro Gln Glr Phe His Asp Ser 105	Ser I Asn S Asn I Leu S Gln G	His Leu Ser Ala Pro Gly 60 Ser Glu 75 Arg Arg	Pro G Gly G 45 Pro A Gly S Gln G	I5 Gly Ala 30 Gly Pro Arg Ser Ger Met 95 Gln Gln	Asp Thr Lys Phe 80 Glu Gln

Met His Arg Val Ser Tyr Ala Met Ser Leu His Asp Leu Pro Ala Arg

	130					135					140				
Pro	Thr	Ala	Phe	Asn	Arg	Val	Leu	Gln	Gln	Ile	Arg	Ser	Arg	Pro	Ser
145					150					155					160
He	Lys	Arg	Gly	Ala	Ser	Leu	His	Ser	Ser	Ser	Gly	Gly	Gly	Ser	Ser
				165					170					175	
Gly	Ser	Ser	Ser	Arg	Arg	Thr	Lys	Ser	Ser	Ser	Leu	Glu	Pro	Gln	Arg
			180					185					190		
Gly	Ser	Pro	His	Leu	Leu	Arg	Lys	Ala	Pro	Gln	Asp	Ser	Ser	Leu	Ala
		195					200					205			
Ala	He	Leu	His	Gln	His	Gln	Cys	Arg	Pro	Arg	Ser	Ser	Ser	Thr	Thr
	210					215					220				
Asp	Thr	Ala	Leu	Leu	Leu	Ala	Asp	Gly	Ser	Asn	Val	Tyr	Leu	Leu	Ala
225					230					235					240
G] u	Glu	Ala	Glu	Gly	He	G1 y	Asp	Lys	Val	Asp	Lys	Gly	Asp	Leu	Val
				245					250					255	
Ala	Leu	Ser		Pro	Ala	Gly	His		Asp	Thr	Asp	Gly		lle	Ser
			260					265					270		
Leu	Asp		Pro	Asp	Gly	Ala		Asp	Pro	Gln	Arg		Lys	Ala	Ala
		275					280	_	_			285			
He		His	Leu	His	Gln		He	Leu	Lys	He		Glu	GIn	Лe	Lys
7.1	290	0.1	0.1			295			W 1	4.7	300	T	,	,	,
	Glu	GIn	Glu	Ala	Arg	Asp	Asp	Asn	val		Glu	lyr	Leu	Lys	
305	Δ	Δ	A1.	Λ	310	C1	C1	V - 1	C	315	11.	1	C1	V - 1	320 Db-
АТа	ASII	ASN	Ala	325	Lys	GIII	GIN	vai	330	Arg	116	Lys	GIII	335	rne
C1 ₁₁	Lvc	lve	Acn		Lys	Sor	Λla	Gla		Ho	Δla	Gln	Lou		lve
Olu	Lys	Lys	340	0111	Lys	361	MIG	345		110	MIG	OIII	350	1113	Lys
Lve	Len	Glu		Tvr	Arg	Arø	Arø			Glu	11e	Glu		Asn	Glv
Lys	Lea	355	nis	1 , 2	6	/11 B	360	БСС	2,5	014	110	365	0111	11511	01,
Pro	Ser		Gln	Pro	Lys	Asp		Leu	Arg	Asp	Met		Gln	Gly	Leu
	370	6			_,_	375			0		380				
Lys		Val	Gly	Ala	Asn		Arg	Ala	Gly	lle		Gly	Phe	Gly	Gly
385	•		•		390		J		-	395		-		-	400
	Val	Val	Glu	Gly	Val	Lys	Gly	Ser	Leu	Ser	Gly	Leu	Ser	Gln	Ala
-				405					410					415	
Thr	Hic	Thr	Λla	Val	Val	Sor	Lvc	Pro	Ara	Glu	Pho	Ala	Ser	Len	He

			420					425					430		
Arg	Asn	Lys	Phe	Gly	Ser	Ala	Asp	Asn	He	Ala	His	Leu	Lys	Asp	Pro
		435					440					445			
Leu	Glu	Asp	Gly	Pro	Pro	Glu	Glu	Ala	Ala	Arg	Ala	Leu	Ser	Gly	Ser
	450					455					460				
Ala	Thr	Leu	Val	Ser	Ser	Pro	Lys	Tyr	Gly	Ser	Asp	Asp	Glu	Cys	Ser
465					470					475					480
Ser	Ala	Ser	Ala	Ser	Ser	Ala	Gly	Ala	Gly	Ser	Asn	Ser	Gly	Ala	Gly
				485					490					495	
Pro	Gly	Gly	Ala	Leu	Gly	Ser	Pro	Lys	Ser	Asn	Ala	Leu	Tyr	Gly	Ala
			500					505					510		
Pro	Gly	Asn	Leu	Asp	Ala	Leu	Leu	Glu	Glu	Leu	Arg	Glu	Ile	Lys	Glu
		515					520					525			
Gly	Gln	Ser	His	Leu	Glu	Asp	Ser	Met	Glu	Asp	Leu	Lys	Thr	Gln	Leu
	530					535					540				
Gln	Arg	Asp	Tyr	Thr	Tyr	Met	Thr	Gln	Cys	Leu	Gln	Glu	Glu	Arg	Tyr
545					550					555					560
Arg	Tyr	Glu	Arg	Leu	Glu	Glu	Gln	Leu	Asn	Asp	Leu	Thr	Glu	Leu	His
				565					570					575	
Gln	Asn	Glu	Met	Thr	Asn	Leu	Lys	Gln	Glu	Leu	Ala	Ser	Met	Glu	Glu
			580					585					590		
Lys	Val	Ala	Tyr	Gln	Ser	Tyr	Glu	Arg	Ala	Arg	Asp	Ile	Gln	Glu	Ala
		595					600					605			
Val	${\tt Glu}$	Ser	Cys	Leu	Thr	Arg	Val	Thr	Lys	Leu	Glu	Leu	Gln	Gln	Gln
	610					615					620				
Gln	Gln	Gln	Val	Val	Gln	Leu	Glu	Gly	Val	Glu	Asn	Ala	Asn	Ala	Arg
625					630					635					640
Ala	Leu	Leu	Gly	Lys	Phe	Ile	Asn	Val	lle	Leu	Ala	Leu	Met	Ala	Val
				645					650					655	
Leu	Leu	Val	Phe	Val	Ser	Thr	lle	Ala	Λsn	Phe	lle	Thr	Pro	Leu	Met
			660					665					670		
Lys	Thr	Arg	Leu	Arg	lle	Thr	Ser	Thr	Thr	Leu	Leu	Val	Leu	Val	Leu
		675					680					685			
Phe	Leu	Leu	Trp	Lys	His	Trp	Asp	Ser	Leu	Thr	Tyr	Leu	Leu	Glu	His
	690					695					700				
Va]	Leu	Leu	Pro	Ser											

705

<210> 4842

<211> 335 <212> PRT <213> Homo sapiens <400> 465 Met Gln Arg His Val Met Val Ala Asn Pro Ser Val Leu Cys Ser His 10 Phe Asn Gln Asp Leu Trp Pro Glu Gln Ser Ile Lys Asp Ser Phe Gln 25 Lys Leu 11e Leu Arg Arg His Lys Lys Cys Gly His Asp Asn Leu Gln 40 Leu Lys Lys Gly Cys Glu Ser Val Asp Lys Cys Lys Val His Lys Arg 55 60 Gly Tyr Asn Gly Leu Asn Gln Cys Leu Thr Thr Gln Ser Lys Met 70 65 75 80 Phe Gln Cys Asp Lys His Gly Lys Val Phe His Gln Phe Ser Asn Thr 90 Asn Arg His Lys Ile Arg His Thr Gly Lys Asn Pro Cys Lys Phe Thr 105 Glu Cys Gly Lys Ala Phe Asn Arg Ser Ser Thr Phe Thr Thr His Lys 115 120 Lys lle His Thr Gly Glu Lys Pro Tyr Lys Cys lle Glu Cys Gly Lys 135 Ala Phe Asn Arg Ser Ser His Leu Thr Thr His Lys lle lle His Thr 145 160 150 155 Gly Glu Lys Arg Tyr Lys Cys Glu Asp Cys Gly Lys Ala Phe Asn Arg 170 Ser Ser Asn Leu Thr Thr His Lys Lys Ile His Thr Gly Glu Lys Pro 180 185 190 Tyr Lys Cys Glu Glu Cys Gly Lys Ala Phe Lys Arg Ser Ser lle Leu 195 200 205

Thr Thr His Lys Arg lle His Thr Gly Glu Lys Pro Tyr Lys Cys Glu

Glu Cys Gly Lys Val Phe Lys Tyr Leu Ser Ser Leu Ser Thr His Lys Ile Ile His Thr Gly Glu Lys Pro Tyr Lys Cys Glu Glu Cys Gly Lys Ala Phe Asn Trp Ser Ser His Leu Thr Thr His Lys Arg Ile His Thr Gly Glu Lys Pro Tyr Lys Cys Glu Glu Cys Gly Lys Gly Phe Lys Tyr Ser Ser Thr Leu Thr Lys His Lys Ile Ile His Thr Gly Glu Lys Pro Tyr Lys Cys Glu Glu Cys Arg Ser Leu Arg Ser Gln Cys Asp Gln Leu Glu Glu Arg Val Ser Val Met Glu Asp Glu Met Asn Gly Met Lys

<210> 4843

<211> 505

<212> PRT

<213> Homo sapiens

<400> 466

Met Lys Ala Gln Arg Glu Arg Leu Gln Ile Pro Gly Leu Thr Leu Asp Leu Thr Pro Arg Ser Leu Ser Pro Thr Pro Ser Ser Pro Gly Ser Pro Cys Ser Pro Leu Leu Ala Phe His Phe Trp Ser Cys Arg Thr Ser Asn Arg Lys Ser Leu lle Gly Asn Gly Gln Scr Pro Ala Leu Pro Arg Pro His Ser Pro Leu Ser Ala His Ala Gly Asn Ser Pro Gln Asp Ser Pro Arg Asn Phe Ser Pro Ser Ala Ser Ala His Phe Ser Phe Ala Arg Arg

Asn Asp Arg Thr Asp Gly Arg Arg Trp Ser Leu Ala Ser Leu Pro Ser

			100					105					110		
Ser	Gly	Tyr	Gly	Thr	Asn	Thr	Pro	Ser	Ser	Thr	Val	Ser	Ser	Ser	Cys
		115					120					125			
Ser	Ser	Gln	Glu	Lys	Leu	His	Gln	Leu	Pro	Tyr	Gln	Pro	Thr	Pro	Asp
	130					135					140				
Glu	Leu	His	Phe	Leu	Ser	Lys	His	Phe	Cys	Thr	Thr	Glu	Ser	lle	Ala
145					150					155					160
Thr	Glu	Asn	Arg	Cys	Arg	Asn	Thr	Pro	Met	His	Pro	Arg	Ser	Arg	Ser
				165					170					175	
Leu	Ser	Pro	Gly	Arg	Ser	Pro	Ala	Cys	Cys	Asp	His	Glu	He	He	Met
			180					185					190		
Met	Asn	His	Val	Tyr	Lys	Glu	Arg	Phe	Pro	Lys	Ala	Thr	Ala	Gln	Met
		195					200					205			
Glu	Glu	Arg	Leu	Lys	Glu	Ile	He	Thr	Ser	Tyr	Ser	Pro	Asp	Asn	Val
	210					215					220				
Leu	Pro	Leu	Ala	Asp	Gly	Val	Leu	Ser	Phe	Thr	His	His	Gln	Ile	He
225					230					235					240
Glu	Leu	Ala	Arg	Asp	Cys	Leu	Asp	Lys	Ser	His	Gln	Gly	Leu	Ile	Thr
				245					250					255	
Ser	Arg	Tyr	Phe	Leu	Glu	Leu	Gln	His	Lys	Leu	Asp	Lys	Leu	Leu	Gln
			260					265					270		
Glu	Ala	His	Asp	Arg	Ser	Glu	Ser	Gly	Glu	Leu	Ala	Phe	lle	Lys	G1n
		275					280					285			
Leu	Val	Arg	Lys	He	Leu	Ile	Val	lle	Ala	Arg	Pro	Ala	Arg	Leu	Leu
	290					295					300				
Glu	Cys	Leu	Glu	Phe	Asp	Pro	Glu	Glu	Phe	Tyr	Tyr	Leu	Leu	Glu	Лlа
305					310					315					320
Ala	Glu	Gly	His	Ala	Lys	Glu	Gly	Gln		He	Lys	Thr	Asp		Pro
				325					330					335	
Arg	Tyr	lle	lle	Ser	Gln	Leu	Gly	Leu	Asn	Lys	Asp	Pro	Leu	Glu	Glu
			340					345					350		
Met	Ala	His	Leu	G1y	Asn	Tyr		Ser	G1 y	Thr	Ala		Thr	Pro	G] u
		355					360					365			
Thr		Glu	Ser	Val	Ser		Ser	Asn	Ala	Ser		Lys	Leu	Arg	Arg
	370					375				_	380				
lve	Pro	Arg	Glu	Ser	Asp	Phe	Glu	Thr	He	LVS	PH	He	Ser	Asn	Glv

385					390					395					400
Ala	Tyr	Gly	Ala	Val	Tyr	Phe	Val	Arg	His	Lys	Glu	Ser	Arg	Gln	Arg
				405					410					415	
Phe	Ala	Met	Lys	Lys	He	Asn	Lys	Gln	Asn	Leu	Пе	Leu	Arg	Asn	Gln
			420					425					430		
He	Gln	Gln	Ala	Phe	Val	Glu	Arg	Asp	lle	Leu	Thr	Phe	Ala	Glu	Asn
		435					440					445			
Pro	Phe	Val	Val	Ser	Met	Tyr	Cys	Ser	Phe	Glu	Thr	Arg	Arg	His	Leu
	450					455					460				
Cys	Met	Val	Met	Glu	Tyr	Val	Glu	Gly	Gly	Asp	Cys	Ala	Thr	Leu	Met
465					470					475					480
Lys	Asn	Thr	Gly	Pro	Leu	Pro	Val	Asp	Met	Ala	Arg	Met	Tyr	Phe	Ala
				485					490					495	
Glu	Thr	Val	Leu	Ala	Leu	Glu	Phe	Thr							
			500					505							
<210)> 48	344													
<21	1> 40	80													
<212	2> PI	RT													
<213	3> He	omo :	sapi	ens											

Met Arg Glu Lys Ser Phe Gln Cys Asn Glu Ser Gly Lys Ala Phe Asn Tyr Ser Ser Leu Leu Arg Lys His Gln Ile lle His Leu Gly Glu Lys Gln Tyr Lys Cys Asp Val Cys Gly Lys Val Phe Asn Arg Lys Arg Asn Leu Val Cys His Arg Arg Cys His Thr Gly Glu Lys Pro Tyr Arg Cys Asn Glu Cys Gly Lys Thr Phe Ser Gln Thr Tyr Ser Leu Thr Cys His Arg Arg Leu His Thr Gly Glu Lys Pro Tyr Lys Cys Glu Glu Cys Asp

<400> 467

Lys	Ala	Phe	Ser 100	Phe	Lys	Ser	Asn	Leu 105	Lys	Arg	His	Arg	Arg 110	lle	His
Λla	C1	C1		Pro	Tur	Lya	Cva		Clu	Cvc	Cl _v	Lvc		Dho	Sor
ліа	Gly	115	Lys	110	1 y 1	Lys	120	ASII	Olu	Cys	Uly	125	1111	1 116	361
C15	Thu		Com	Lau	The	Cua		A 22.07	A 22 cr	Lou	ui e		C1 _v	C1	Luc
GIII	130	361	261	Leu	1111	135	1115	W1 B	nı g	Leu	140	1111	Gry	Olu	Lys
Duo		Luc	Cua	Asn	C1		C1	Luc	The	Dho		A 25.07	Luc	Sor	Sor
	rne	LyS	Cys	ASII	150	Cys	Gry	Lys	1111	155	361	Mg	Lys	261	
145	The	Cvo	ui o	His		Lou	ui o	The	C1		Lvc	Dro	Tur	Lvo	160
Leu	1111	Cys	1115	165	AI g	Leu	1115	1111	170	Olu	Lys	110	1 9 1	175	Cys
Acn	C1u	Cvc	Cly		Thr	Dho	Sor	Cln.		Lou	Thr	Lou	Lvc		Hic
ASII	Giu	Cys	180	Lys	1111	rne	261	185	Gju	Leu	1111	Leu	190	Cys	1112
Ara	Ara	Lau		Thr	Clv	Clu	Lvc		Tyr	lvc	Cvc	Acn		Cvc	Clv
ΛI g	Mg	195	1115	1 1111	OTY	Olu	200	110	1 y 1	ris	Cys	205	Oju	Cys	Gry
Lvc	Val		Acn	Lys	Lvc	Ala		Lou	Δ1a	Ara	Hic		Δησ	Lou	Hic
rys	210	THE	11311	Lys	1. y 3	215	ASII	Leu	MIG	ni g	220	1113	M B	LCu	1113
Ser		Glu	lve	Pro	Tyr		Cvs	Thr	Glu	Cvs		Lvs	Thr	Phe	Ser
225	01)	oru	12,5	110	230	2,5	0,5		014	235	741	Lys	7111	1110	240
	Asn	Ser	Ala	Leu		He	His	Lvs	Ala		His	He	Glv	Glu	
0				245				, -	250				,	255	-3-
Arg	Tyr	Lvs	Cys	Asn	Glu	Cys	Gly	Lys		Phe	Ser	Arg	lle		Ala
			260					265					270		
Leu	Val	Пe	llis	Thr	Ala	lle	His	Thr	Gly	Glu	Lys	Pro	Tyr	Lys	Cys
		275					280					285			
Asn	Glu	Cys	Gly	Lys	Gly	Phe	Asn	Arg	Lys	Thr	His	Leu	Ala	Cys	His
	290					295					300				
His	Arg	Leu	llis	Thr	Gly	Glu	Lys	Pro	Tyr	Lys	Cys	Asn	Glu	Cys	Gly
305					310					315					320
Lys	Val	Phe	Asn	Arg	Lys	Thr	His	Leu	Ala	His	His	His	Arg	Leu	His
				325					330					335	
Thr	Gly	Asp	Lys	Pro	Tyr	Lys	Cys	Asn	Ġlu	Cys	Gly	Lys	Val	Phe	Asn
			340					345					350		
Gln	Lys	Ala	His	Leu	Ala	Arg	His	His	Arg	Leu	His	Thr	Gly	G] u	Lys
		355					360					365			
Pro	Tyr	Lys	Cys	Asn	Glu	Cys	Gly	Lys	Va]	Phe	Asn	Gln	Lys	Ala	Asn
	370					375					380				

Leu Ala Arg His His Arg Leu His Thr Gly Glu Lys Pro Tyr Lys Phe
385 390 395 400

Asn Glu Cys Gly Lys Ala Phe Asn
405

<210> 4845

<211> 596

<212> PRT

<213> Homo sapiens

<400> 468

Met Leu Pro Ser Gln Ala Gln Gly Leu Leu Trp Trp Val Phe Pro Leu

1 5 10 15

Phe Pro Ala Ser Ser Leu Ser Tyr Pro Pro Val Ser Tyr Arg Ala Asp
20 25 30

Gly Leu Ala Arg Asn Thr Phe Leu Lys Ala Cys Ser Glu Ser Ser Ser 35 40 45

Ser Ser Asn Ile Ser Thr Met Leu Val Thr His Asp Tyr Thr Ala Val 50 55 60

Lys Glu Asp Glu Ile Asn Val Tyr Gln Gly Glu Val Val Gln Ile Leu 65 70 75 80

Ala Ser Asn Gln Gln Asn Met Phe Leu Val Phe Arg Ala Ala Thr Asp 85 90 95

Gln Cys Pro Ala Ala Glu Gly Trp Ile Pro Gly Phe Val Leu Gly His 100 105 110

Thr Ser Ala Val Ile Val Glu Asn Pro Asp Gly Thr Leu Lys Lys Ser 115 120 125

Thr Ser Trp His Thr Ala Leu Arg Leu Arg Lys Lys Ser Glu Lys Lys 130 135 140

Asp Lys Asp Gly Lys Arg Glu Gly Lys Leu Glu Asn Gly Tyr Arg Lys 145 150 155 160

Ser Arg Glu Gly Leu Ser Asn Lys Val Ser Val Lys Leu Leu Asn Pro 165 170 175

Asn Tyr Ile Tyr Asp Val Pro Pro Glu Phe Val Ile Pro Leu Ser Glu 180 185 190

Val	Thr	Cys	Glu	Thr	Gly	Glu	Thr	Val	Val	Leu	Arg	Cys	Arg	Val	Cys
		195					200					205			
Gly	Arg	Pro	Lys	Ala	Ser	He	Thr	Trp	Lys	Gly	Pro	Glu	His	Asn	Thr
	210					215					220				
Leu	Asn	Asn	Asp	Gly	His	Tyr	Ser	Пе	Ser	Tyr	Ser	Asp	Leu	Gly	Glu
225					230					235					240
Ala	Thr	Leu	Lys	He	Val	G] y	Val	Thr	Thr	Glu	Asp	Asp	Gly	Пе	Tyr
				245					250					255	
Thr	Cys	Ile	Ala	Val	Asn	Asp	Met	Gly	Ser	Ala	Ser	Ser	Ser	Ala	Ser
			260					265					270		
Leu	Arg	Val	Leu	Gly	Pro	Gly	Met	Asp	Gly	He	Met	Val	Thr	Trp	Lys
		275					280					285			
Asp	Asn	Phe	Asp	Ser	Phe	Tyr	Ser	G]u	Val	Ala	Glu	Leu	Gly	Arg	Gly
	290					295					300				
Arg	Phe	Ser	Val	Val	Lys	Lys	Cys	Asp	Gln	Lys	Gly	Thr	Lys	Arg	Ala
305					310					315					320
Val	Ala	Thr	Lys	Phe	Va]	Asn	Lys	Lys	Leu	Met	Lys	Arg	Asp	Gln	Val
				325					330					335	
Thr	His	Glu	Leu	Gly	lle	Leu	Gln	Ser	Leu	Gln	His	Pro	Leu	Leu	Val
			340					345					350		
Gly	Leu	Leu	Asp	Thr	Phe	Glu	Thr	Pro	Thr	Ser	Tyr	He	Leu	Val	Leu
		355					360					365			
Glu	Met	Ala	Asp	Gln	Gly	Arg	Leu	Leu	Asp	Cys	Val	Va]	Arg	Trp	Gly
	370					375					380				
Ser	Leu	Thr	Glu	Gly	Lys	11e	Arg	Ala	His	Leu	Gly	Glu	Val	Leu	Glu
385					390					395					400
Ala	Val	Arg	Tyr	Leu	His	Asn	Cys	Arg	11e	Ala	His	Leu	Asp	Leu	Lys
				405					410					415	
Pro	G]u	Asn	lle	Leu	Val	Asp	Glu	Ser	Leu	Ala	Lys	Pro	Thr	He	Lys
			420					425					430		
Leu	Ala	Asp	Phe	Gly	Asp	Ala	Va]	Gln	Leu	Asn	Thr	Thr	Tyr	Tyr	11e
		435					440					445			
His	Gln	Leu	Leu	Gly	Asn	Pro	Glu	Phe	Ala	Ala	Pro	Glu	Ile	11e	Leu
	450					455					460				
Gly	Λsn	Pro	Val	Ser	Leu	Thr	Ser	Asp	Thr	Trp	Ser	Val	Gly	Val	Leu

Thr Tyr Val Leu Leu Ser Gly Val Ser Pro Phe Leu Asp Asp Ser Val Glu Glu Thr Cys Leu Asn Ile Cys Arg Leu Asp Phe Ser Phe Pro Asp Asp Tyr Phe Lys Gly Val Ser Gln Lys Ala Lys Glu Phe Val Cys Phe Leu Leu Gln Glu Asp Pro Ala Lys Arg Pro Ser Ala Ala Leu Ala Leu Gln Glu Gln Trp Leu Gln Ala Gly Asn Gly Arg Ser Thr Gly Val Leu Asp Thr Ser Arg Leu Thr Ser Phe lle Glu Arg Arg Lys His Gln Asn Asp Val Arg Pro Ile Arg Ser Ile Lys Asn Phe Leu Gln Ser Arg Leu Leu Pro Arg Val

<210> 4846

<211> 569

<212> PRT

<213> Homo sapiens

<400> 469

Met Pro Ser Ser Leu Phe Ala Asp Leu Glu Arg Asn Gly Ser Gly Gly Gly Gly Gly Gly Ser Ser Gly Gly Gly Glu Thr Leu Asp Asp Gln Arg Ala Leu Gln Leu Ala Leu Asp Gln Leu Ser Leu Leu Gly Leu Asp Ser Asp Glu Gly Ala Ser Leu Tyr Asp Ser Glu Pro Arg Lys Lys Ser Val Asn Met Thr Glu Cys Val Pro Val Pro Ser Ser Glu His Val Ala Glu lle Val Gly Arg Gln Gly Cys Lys lle Lys Ala Leu Arg Ala Lys Thr

				85					90					95	
Asn	Thr	Tyr	lle	Lys	Thr	Pro	Val	Arg	Gly	Glu	Glu	Pro	Val	Phe	Val
			100					105					110		
Val	Thr	Gly	Arg	Lys	Glu	Asp	Val	Ala	Met	Ala	Arg	Arg	Glu	lle	Пe
		115					120					125			
Ser	Ala	Ala	Glu	His	Phe	Ser	Met	He	Arg	Ala	Ser	Arg	Asn	Lys	Asn
	130					135					140				
Thr	Ala	Leu	Asn	Gly	Ala	Val	Pro	G1 y	Pro	Pro	Asn	Leu	Pro	Gly	Gln
145					150					155					160
Thr	Thr	Ile	Gln	Val	Arg	Val	Pro	Tyr	Arg	Val	Val	Gly	Leu	Val	Val
				165					170					175	
Gly	Pro	Lys	Gly	Ala	Thr	lle	Lys	Arg	Лlе	Gln	Gln	Gln	Thr	His	Thr
			180					185					190		
Tyr	lle	Va]	Thr	Pro	Ser	Arg	Лѕр	Lys	Glu	Pro	Val	Phe	Glu	Val	Thr
		195					200					205			
Gly	Met	Pro	Glu	Asn	Val	Asp	Arg	Ala	Arg	Glu	Glu	He	Glu	Ala	His
	210					215					220				
Ile	Ala	Leu	Arg	Thr	Gly	Gly	lle	He	Glu		Thr	Asp	Glu	Asn	Asp
225					230					235					240
Phe	His	Ala	Asn	Gly	Thr	Asp	Val	Gly		Asp	Leu	His	His		Ser
				245					250					255	
Gly	Gly	Ser		Pro	Gly	Ser	Leu		Ser	Lys	Pro	Thr	Pro	Ser	He
			260					265					270		
Thr	Pro		Pro	Gly	Arg	Lys		Phe	Ser	Ser	lyr		Asn	Asp	Ser
	C	275		61	C		280	TI		C	т.	285	C1	C1	C1
Ser			Leu	61 y	Ser			Inr	Asp	Ser			Gly	GIŅ	GIŸ
Tl	290		C a se	۸1.	110	295		Cln	A 22.00	Lau	300		Tue	Sor	Dro
	ser	ser	ser	MIa		Ala	1111	GIII	AIG	315	на	ush	Tyr	261	320
305	Sor	Dro	410	Lou	310	Pho	Λla	Hic	Acn		Aen	Aen	Asn	Aen	
F10	561	110	Міа	325	561	1116	ЛΙα	1112	330	Oly	nan		или	335	11-311
Glv	Asn	Glv	Tyr		Tyr	Thr	Ala	Glv		Glu	Ala	Ser	Val		Ser
Gry	11311	01,	340	1 . 1.1	1 7 3	111.	7,74	345	01)	014			350		0
Pro	Asn	Glv		Pro	Glu	Leu	Gln		Thr	Phe	Asp	Pro	Ala	Pro	Ala
.10		355					360				I-	365			
Pro	Pro			Ala	Pro	Leu		Trp	Ala	Gln	Phe		Arg	Ser	Pro

Gly Gly Gly Pro Ala Ala Pro Val Ser Ser Ser Cys Ser Ser Ser Ala Ser Ser Ser Ala Ser Ser Ser Ser Val Val Phe Pro Gly Gly Ala Ser Ala Pro Ser Asn Ala Asn Leu Gly Leu Leu Val His Arg Arg Leu His Pro Gly Thr Ser Cys Pro Arg Leu Ser Pro Pro Leu His Met Ala Pro Gly Ala Gly Glu His His Leu Ala Arg Arg Val Arg Ser Asp Pro Gly Gly Gly Leu Ala Tyr Ala Ala Tyr Ala Asn Gly Leu Gly Ala Gln Leu Pro Gly Leu Gln Pro Ser Asp Thr Ser Gly Ser Ser Ser Lys Gly Ser Arg Asp Cys Ser Val Cys Phe Glu Ser Glu Val Ile Ala Ala Leu Val Pro Cys Gly His Asn Leu Phe Cys Met Glu Cys Ala Asn Arg Ile Cys Glu Lys Ser Glu Pro Glu Cys Pro Val Cys His Thr Ala Val Thr Gln Ala Ile Arg Ile Phe Ser

<210> 4847

<211> 302

<212> PRT

<213> Homo sapiens

<400> 470

Met Ala Leu Leu Ile Thr Pro Ala Gly Val Ala Thr Val Asn Arg His

1 5 10 15

Ser Thr Ile Pro Ser Asp Thr His Thr Ser Arg Glu Lys Pro Arg Phe

			20					25					30		
His	Lys	Pro	Cys	Arg	Asn	Asp	Leu	Glu	Ser	Leu	Leu	Ser	Glu	Gly	Arg
		35					40				٠	45			
Leu	Asp	Thr	Ser	Val	Gln	Thr	Pro	Cys	Pro	Gln	His	Pro	His	Thr	Gln
	50					55					60				
Leu	Ser	Cys	Glu	Pro	Gln	Pro	Leu	Glu	His	Ser	Ser	Cys	Leu	Ser	Thr
65					70					75					80
Cys	Leu	Ala	Gly	Cys	Phe	Leu	Pro	Val	Pro	Ser	Ser	Pro	His	Thr	His
				85					90					95	
Pro	Leu	Leu	Pro	Gly	Ser	Arg	Trp	Leu	Pro	Pro	Pro	Leu	Ala	Leu	Leu
			100					105					110		
Met	Gly	Thr	Leu	Ser	Pro	Gly	Leu	Ala	Val	Lys	Pro	Ser	Trp	Val	Pro
		115					120					125			
Arg	Phe	Pro	Leu	Leu	Ala	Arg	Gln	Ser	Pro	Ala	Thr	Ser	Va]	Gly	Met
	130					135					140				
Pro	Leu	Ser	Ala	Ala		Gln	Pro	Gly	Ser		Gly	Arg	Leu	His	
145					150					155					160
Pro	Lys	Leu	Arg		Ser	Ser	Pro	Phe		Gly	His	Ser	Asp		Asn
				165					170			_		175	_
Lys	Ala	Thr		Gln	Gly	Arg	Glu		Arg	Asp	GIn	Pro		Arg	Pro
			180	0.1	0	Б	0.1	185			0.1	6	190	m.	
Ser	His	Leu	Cys	Glu	Cys	Pro		Ala	Ala	Lys	GIn		Ala	Hhr	Asn
C1	V - 1	195	C1	T1	Λ	Λ	200	M - 1	DL.	D	1	205	C	C1	11
GIY		Ala	GIU	ınr	ASN		Ser	vai	Pne	Pro		GTY	ser	GJU	мла
Ana	210	Leu	Sar	Lou	Ara	215	Cln	Clu	Sor	Cln	220 Pro	Hic	Sor	C1v	Sor
225	361	Leu	361	Leu	230		GIII	01u	261	235	110	1112	361	ОТУ	240
	Δra	Arg	Gla	Sor			Cve	Sor	Pro		Pho	Trn	Cve	Cve	
561	ni g	Mi g	Olu	245	, (11	561	Cys	561	250	261	1116	цр	Cys	255	пр
Gln	Pro	Leu	Ala		Leu	Thr	Cvs	G1 v		Ala	Ala	Pro	He		Val
OIII		LCG	260	1	Lou	7117	0,5	265	0,5	711 (7,10	110	270	501	,
Pro	Glv	Val		Arg	Pro	Ser	Pro		Pro	Cvs	Cvs	Val		Pro	Pro
	- 3	275					280	3		• -		285		-	-
Leu	Val	Arg	Leu	Gln	Ser	Leu		Leu	Gly	Pro	Thr		He		
	290	9				295	•		·		300				

```
<211> 449
<212> PRT
<213> Homo sapiens
<400> 471
Met Pro Gly Met Met Glu Lys Gly Pro Glu Leu Leu Gly Lys Asn Arg
 1
                  5
                                     10
                                                          15
Ser Ala Asn Gly Ser Ala Lys Ser Pro Ala Gly Gly Gly Ser Gly
                                 25
Ala Ser Ser Thr Asn Gly Gly Leu His Tyr Ser Glu Pro Glu Ser Gly
                             40
                                                 45
Cys Ser Ser Asp Asp Glu His Asp Val Gly Met Arg Val Gly Ala Glu
     50
                         55
                                             60
Tyr Gln Ala Arg Ile Pro Glu Phe Asp Pro Gly Ala Thr Lys Tyr Thr
                     70
                                         75
Asp Lys Asp Asn Gly Gly Met Leu Val Trp Ser Pro Tyr His Ser Ile
                 85
                                     90
Pro Asp Ala Lys Leu Asp Glu Tyr Ile Ala Ile Ala Lys Glu Lys His
            100
                                105
                                                     110
Gly Tyr Asn Val Glu Gln Ala Leu Gly Met Leu Phe Trp His Lys His
                            120
                                                 125
Asn Ile Glu Lys Ser Leu Ala Asp Leu Pro Asn Phe Thr Pro Phe Pro
    130
                        135
                                             140
Asp Glu Trp Thr Val Glu Asp Lys Val Leu Phe Glu Gln Ala Phe Ser
                    150
                                        155
Phe His Gly Lys Ser Phe His Arg Ile Gln Gln Met Leu Pro Asp Lys
                165
                                    170
                                                         175
Thr lle Ala Ser Leu Val Lys Tyr Tyr Tyr Ser Trp Lys Lys Thr Arg
            180
                                185
Ser Arg Thr Ser Leu Met Asp Arg Gln Ala Arg Lys Leu Ala Asn Arg
                            200
                                                 205
His Asn Gln Gly Asp Ser Asp Asp Val Glu Glu Thr His Pro Met
    210
                        215
                                             220
```

Asp Gly Asn Asp Ser Asp Tyr Asp Pro Lys Lys Glu Ala Lys Lys Glu

<210> 4848

225					230					235					240
Gly	Asn	Thr	Glu	Gln	Pro	Val	Gln	Thr	Ser	Lys	Пe	Gly	Leu	G1 y	Arg
				245					250					255	
Arg	Glu	Tyr	Gln	Ser	Leu	Gln	His	Arg	His	His	Ser	Gln	Arg	Ser	Lys
			260					265					270		
Cys	Arg	Pro	Pro	Lys	Gly	Met	Tyr	Leu	Thr	Gln	Glu	Asp	Val	Val	Ala
		275					280					285			
Val	Ser	Cys	Ser	Pro	Asn	Ala	Ala	Asn	Thr	He	Leu	Arg	Gln	Leu	Asp
	290					295					300				
Met	Glu	Leu	He	Ser	Leu	Lys	Arg	Gln	Val	Gln	Asn	Ala	Lys	Gln	Val
305					310					315					320
Asn	Ser	Ala	Leu	Lys	Gln	Lys	Met	Glu	Gly	Gly	He	Glu	Glu	Phe	Lys
				325					330					335	
Pro	Pro	Glu	Ser	Asn	Gln	Lys	He	Asn	Ala	Arg	Trp	Thr	Thr	Glu	Glu
			340					345					350		
Gln	Leu	Leu	Ala	Va]	Gln	Gly	Val	Arg	Lys	Tyr	Gly	Lys	Asp	Phe	Gln
		355					360					365			
Ala	He	Ala	Asp	`Val	Ile	Gly	Asn	Lys	Thr	Val	Gly	Gln	Val	Lys	Asn
	370					375					380				
Phe	Phe	Va]	Asn	Tyr	Arg	Arg	Arg	Phe	Asn	Leu	Glu	Glu	Val	Leu	Gln
385					390					395					400
Glu	Trp	Glu	Ala	Glu	Gln	Gly	Thr	Gln	Ala	Ser	Asn	Gly	Asp	Ala	Ser
				405					410					415	
Thr	Leu	Gly	Glu	Glu	Thr	Lys	Ser	Ala	Ser	Asn	Val	Pro	Ser	Gly	Lys
			420					425					430		
Ser	Thr	Asp	Glu	Glu	Glu	Glu	Val	Cys	Leu	Cys	Met	Glu	Phe	Glu	Leu
		435					440					445			
lle															

<210> 4849

<211> 244

<212> PRT

<213> Homo sapiens

<400)> 47	72													
Met	Thr	Gly	Ser	Asn	Ser	His	Ile	Thr	lle	Leu	Thr	Val	Asn	lle	Asn
1				5					10					15	
Gly	Leu	Asn	Ala	Pro	Пе	Lys	Gly	His	Arg	Leu	Ala	Asn	Trp	He	Lys
			20					25					30		
Ser	Gln	Asp	Pro	Ser	Val	Cys	Cys	Ile	Gln	Glu	Thr	His	Leu	Met	Arg
		35					40					45			
Lys	Phe	Thr	His	Arg	Phe	Asn	Ile	Lys	Gly	Trp	Arg	Lys	Ile	Tyr	Gln
	50					55					60				
Glu	Asn	Gly	Lys	Gln	Lys	Lys	Ala	G1 y	Val	Ala	Ile	Leu	Val	Ser	Asp
65					70					7 5					80
Lys	Thr	Asp	Phe	Lys	Pro	Thr	Lys	Ile	Lys	Gly	Asp	Lys	Gly	His	Tyr
				85					90					95	
lle	Met	Val	Lys	Gly	Ser	Met	Gln	Gln	Glu	Glu	Leu	Thr	He	Leu	Asn
			100					105					110		
Ile	Tyr	Ala	Pro	Asn	Thr	Arg	Ala	Pro	Arg	Phe	Ile	Lys	Gln	Val	Leu
		115					120					125			
Arg	Asp	Leu	Gln	Arg	Asp	Leu	Asp	Ser	His	Thr	Ile	Ile	Met	Gly	Asp
	130					135					140				
Phe	Asn	Thr	Glu	Leu	Ser	11e	Leu	Glu	Arg	Ser	Thr	Arg	Gln	Lys	Val
145					150					155					160
Asn	Lys	Asp	lle	Gln	Asp	Leu	Asn	Ser	Ala	Leu	Gln	Gln	Thr	Asp	Pro
				165					170					175	
He	Asp	lle	His	Arg	Asn	Leu	His	Leu	Lys	Ser	Thr	Glu	Tyr	Thr	Leu
			180					185					190		
Phe	Ser	Ala	Pro	His	He	Thr	Tyr	Ser	Lys	Phe	Asp	His	lle	Ile	Gly
		195					200					205			
Ser	Lys	Ala	Leu	Leu	Thr	Lys	Cys	Lys	Arg	Thr	Glu	lle	Thr	Thr	Asr
	210					215					220				
Cys	Leu	Leu	Asp	His	Ser	Ala	lle	Lys	Phe	Glu	Leu	Arg	He	Lys	Lys
225					230					235					240
Leu	Thr	Gln	Asn												

<211> 695

```
<212> PRT
<213> Homo sapiens
<400> 473
Met Val Tyr Pro Tyr Pro Gly Ala Arg Ala Glu Glu Lys Leu Gly Gly
                  5
                                     10
Thr Arg Asp Pro Thr Tyr Gln Glu Arg Ala Ala Pro Gln Thr Gln Pro
             20
                                 25
                                                      30
Leu Gly Lys Glu Thr Asp Ser Leu Ser Ala Gly Phe Val Val Met
Gly Val Asp Leu Ser Arg Cys Gly Pro Asp His Thr Ala Ser Arg Cys
                         55
                                             60
Pro Trp Asp Pro Gly Leu Leu Leu Arg Phe Leu Ala Ala Met Ala Ala
                                         75
 65
                     70
Val Gly Ala Leu Glu Pro Leu Leu Pro Gly Pro Leu Leu Ala Val His
                                     90
                 85
Pro His Ala Gly Thr Ala Pro Pro Ala Asn Gln Leu Pro Trp Pro Val
            100
                                105
                                                     110
Leu Cys Ser Pro Val Ala Gly Val Ile Leu Leu Ala Leu Gly Ala Leu
        115
                            120
                                                 125
Leu Val Leu Gln Leu lle Arg Arg Arg Arg Glu His Gly Ala Leu
                        135
Trp Leu Pro Pro Gly Phe Thr Arg Arg Pro Arg Thr Gln Ser Ala Pro
145
                    150
                                         155
His Arg Arg Arg Pro Pro Leu Gly Glu Asp Ser Ile Gly Leu Lys Ala
                                    170
                165
Leu Lys Pro Lys Ala Glu Val Asp Glu Asp Gly Val Val Met Cys Ser
            180
                                                     190
                                185
Gly Pro Glu Glu Glu Glu Glu Glu Glu Glu Thr Gly Pro Pro Ser Thr
                            200
Cys Gln Leu Trp Ser Leu Ser Gly Gly Cys Gly Ala Leu Pro Gln Ala
                        215
                                             220
Ala Met Leu Thr Pro Pro Gln Glu Ser Glu Met Glu Ala Pro Asp Leu
225
                    230
                                         235
                                                             240
```

Asp Thr Arg Gly Pro Asp Gly Val Thr Pro Leu Met Ser Ala Val Cys

				245					250					255	
Cys	Gly	Glu	Val	Gln	Ser	Gly	Thr	Phe	Gln	G1 y	Ala	Trp	Leu	Gly	Cys
			260					265					270		
Pro	Glu	Pro	Trp	Glu	Pro	Leu	Leu	Asp	Gly	Gly	Ala	Cys	Pro	Gln	Ala
		275					280					285			
His	Thr	Val	Gly	Thr	Gly	Glu	Thr	Pro	Leu	His	Leu	Ala	Ala	Arg	Phe
	290					295					300				
Ser	Arg	Pro	Thr	Ala	Ala	Arg	Arg	Leu	Leu	Glu	Ala	Gly	Ala	Asn	Pro
305					310					315					320
Asn	Gln	Pro	Asp	Arg	Ala	Gly	Arg	Thr	Pro	Leu	His	Ala	Ala	Val	Ala
				325					330					335	
Ala	Asp	Ala	Arg	Glu	Val	Cys	Gln	Leu	Leu	Leu	Arg	Ser	Arg	Gln	Thr
			340					345					350		
Ala	Val	Asp	Ala	Arg	Thr	Glu	Asp	Gly	Thr	Thr	Pro	Leu	Met	Leu	Ala
		355					360					365			
Ala	Arg	Leu	Ala	Val	Glu	Asp	Leu	Val	G] u	Glu	Leu	Ile	Ala	Ala	Gln
	370					375					380				
Ala	Asp	Val	Gly	Ala	Arg	Asp	Lys	Trp	Gly	Lys	Thr	Ala	Leu	His	Trp
385	•				390					395					400
Ala	Ala	Ala	Val	Asn	Λsn	Ala	Arg	Ala	Ala	Arg	Ser	Leu	Leu	Gln	Ala
				405					410					415	
Gly	Ala	Asp	Lys	Asp	Ala	Gln	Asp	Asn	Arg	Glu	Gln	Thr	Pro	Leu	Phe
			420					425					430		
Leu	Ala	Ala	Arg	Glu	Gly	Ala	Val	Glu	Va1	Ala	Gln	Leu	Leu	Leu	Gly
		435					440					445			
Leu	Gly	Ala	Ala	Arg	Glu	Leu	Arg	Asp	Gln	Ala	Gly	Leu	Ala	Pro	Ala
	450					455					460				
Asp	Va]	Ala	His	Gln	Arg	Asn	His	Trp	Asp	Leu	Leu	Thr	Leu	Leu	Glu
465					470					475					480
Gly	Ala	Gly	Pro	Pro	Glu	Ala	Arg	His	Lys	Ala	Thr	Pro	Gly	Arg	Glu
				485					490					495	
Ala	Gly	Pro	Phe	Pro	Arg	Ala	Arg	Thr	Val	Ser	Val	Ser	Va]	Pro	Pro
			500					505					510		
His	Gly	G1 y	Gly	Ala	Leu	Pro	Arg	Cys	Arg	Thr	Leu	Ser	Ala	Gly	Ala
		515					520					525			
Glv	Pro	Arg	Gly	Gly	Gly	Ala	Cys	Leu	Gln	Ala	Arg	Thr	Trp	Ser	Val

	530					535					540				
Asp	Leu	Ala	Ala	Arg	Gly	Gly	Gly	Ala	Tyr	Ser	His	Cys	Arg	Ser	Leu
545					550					555					560
Ser	G1 y	Val	Gly	Ala	Gly	Gly	Gly	Pro	Thr	Pro	Arg	G]y	Arg	Arg	Phe
				565					570					575	
Ser	Ala	Gly	Met	Arg	Gly	Pro	Arg	Pro	Asn	Pro	Ala	lle	Met	Arg	Gly
			580					585					590		
Arg	Tyr	Gly	Val	Ala	Ala	Gly	Arg	Gly	Gly	Arg	Val	Ser	Thr	Asp	Asp
		595					600					605			
Trp	Pro	Cys	Asp	Trp	Val	Ala	Leu	Gly	Ala	Cys	Gly	Ser	Ala	Ser	Asn
	610					615					620				
He	Pro	He	Pro	Pro	Pro	Cys	Leu	Thr	Pro	Ser	Pro	Glu	Arg	Gly	Ser
625					630					635					640
Pro	Gln	Leu	Asp	Cys	Gly	Pro	Pro	Ala	Leu	Gln	Glu	Met	Pro	He	Asn
				645					650					655	
Gln	Gly	Gly	Glu	Gly	Lys	Lys	Ile	Glu	Glu	Tyr	Met	Val	Gly	Arg	Asn
			660					665					670		
Ser	Lys	Asn	Asp	Tyr	Pro	Leu	Lys	Gly	Arg	Leu	Glu	Gly	Leu	Pro	Gly
		675					680					685			
Phe	Lys	Met	Asp	Pro	Pro	Lys									
	690					695									

<210> 4851

<211> 211

<212> PRT

<213> Homo sapiens

<400> 474

 Met
 Ser
 Ser
 Val
 Gly
 Leu
 Phe
 Leu
 Cys
 Phe
 Thr
 Ser
 Cys
 Ser
 Ser
 Ser
 Ser
 Inches
 Ser
 Leu
 Ser
 Leu
 Pro
 Leu
 Pro
 Leu
 Pro
 Ser
 His
 Leu
 Val
 Gly
 Leu
 Leu

 Val
 Ser
 Leu
 Leu
 Asp
 Asp
 Asp
 Pro
 Ala
 Pro
 Ser
 His
 Leu
 Val
 Glu
 Lys
 Ile

 Val
 Tyr
 His
 Ser
 Lys
 Tyr
 Lys
 Pro
 Lys
 Arg
 Leu
 Gly
 Asp
 Ala

60 50 55 Leu Met Lys Leu Ala Gly Pro Leu Thr Phe Asn Glu Met 11e Gln Pro 70 75 65 80 Val Cys Leu Pro Asn Ser Glu Glu Asn Phe Pro Asp Gly Lys Val Cys 90 85 Trp Thr Ser Gly Trp Gly Ala Thr Glu Asp Gly Gly Asp Ala Ser Pro 105 Val Leu Asn His Ala Ala Val Pro Leu Ile Ser Asn Lys Ile Cys Asn 115 120 125 His Arg Asp Val Tyr Gly Gly Ile Ile Ser Pro Ser Met Leu Cys Ala 135 Gly Tyr Leu Thr Gly Gly Val Asp Ser Cys Gln Gly Asp Ser Gly Gly 150 155 Pro Leu Val Cys Gln Glu Arg Arg Leu Trp Lys Leu Val Gly Ala Thr 165 170 Ser Phe Gly Ile Gly Cys Ala Glu Val Asn Lys Pro Gly Val Tyr Thr 180 185 Arg Val Thr Ser Phe Leu Asp Trp Ile His Glu Gln Met Glu Arg Asp 195 200 205 Leu Lys Thr 210 ⟨210⟩ 4852 <211> 193 <212> PRT <213> Homo sapiens <400> 475

 1
 5
 10
 15

 Ala Cys Lys Thr Pro 1le Arg Asn Arg Ala Phe Tyr Met Glu Glu Gly
 20
 25
 30

 Val Pro Tyr Cys Glu Arg Gly Thr His Trp Pro Val Arg Val Arg Arg
 45

 Asp Gly Ala Trp Gly Arg His Glu Ser Arg Ser Ser Phe Ser Leu Pro

Met His Ala Leu Lys Met Thr Trp His Val His Cys Phe Thr Cys Ala

Pro Phe Ser Asp Tyr Glu Lys Met Phe Gly Thr Lys Cys His Gly Cys Asp Phe Lys Ile Asp Ala Gly Asp Arg Phe Leu Glu Ala Leu Gly Phe Ser Trp His Asp Thr Cys Phe Val Cys Ala Val Arg Ala Pro Pro Leu Glu Leu Ser Pro Lys Pro Thr Gly Pro Leu Phe Ile Pro Gln Glu Met Gln Glu Lys Leu Gly Arg Gly Leu Ser Cys Cys Pro Gln Pro His Val Thr Glv Pro Leu Leu Ser Leu Asp Met Ser Asp Gln Pro Gly Arg Lys Asp Leu Leu Gln Glu Gly Gln Ala Ser Leu Gln Glu Pro Cys Leu Leu Ser Cys Val Ser Pro Phe Cys Pro Gln Leu Pro Arg Trp Pro Leu Ala <210> 4853 <211> 134 <212> PRT <213> Homo sapiens <400> 476 Met Cys Gln Val Tyr Phe Val Lys Leu Leu Ile Leu Gly Val Leu Leu Cys Phe Leu Leu Val Arg Gln Asp Leu Thr Leu Ser Pro Arg Leu Glu

Cys Ser Gly Ala Ile Trp Ala His Cys Asn Pro Arg Leu Leu Gly Ile

Ser Asn Leu Pro Ala Leu Ala Ser Gln Val Ala Glu Thr Thr Gly Met

 Ser His His Thr Gln Leu Thr Thr Cys Asn Phe Phe Asp Arg Asp Arg
 Asp Arg Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 Asp Arg
 <210> 4854

<211> 568

<212> PRT

<213> Homo sapiens

<400> 477

130

Met Gly Ser Glu Val Val Ala Gly Asn Ser Val Gly Pro Thr Met Gly 5 10 Ala Ala Ser Ser Gly Pro Leu Pro Pro Pro Pro Pro Pro Leu Pro Pro 25 Ser Ser Asp Thr Pro Glu Thr Val Gln Asn Gly Pro Val Thr Pro Pro 35 40 45 Met Pro Pro Pro Pro Leu Pro Gly Pro Ala Ala Glu Thr Val Pro 55 Ala Pro Pro Leu Ala Pro Pro Leu Pro Ser Ala Pro Pro Leu Pro Gly 70 75 Thr Ser Ser Pro Thr Val Val Phe Asn Ser Gly Leu Ala Ala Val Lys 85 90 95 Ile Lys Lys Pro Ile Lys Thr Lys Phe Arg Met Pro Val Phe Asn Trp 100 105 110 Val Ala Leu Lys Pro Asn Gln lle Asn Gly Thr Val Phe Asn Glu lle 115 120 125

Asp Asp Glu Arg 11e Leu Glu Asp Leu Asn Val Asp Glu Phe Glu Glu

140

He	Phe	Lys	Thr	Lys	Ala	Gln	Gly	Pro	Ala	He	Asp	Leu	Ser	Ser	Ser
145					150					155					160
Lys	Gln	Lys	He	Pro	Gln	Lys	Gly	Ser	Asn	Lys	Val	Thr	Leu	Leu	Glu
				165					170					175	
Ala	Λsn	Arg	Λlа	Lys	Asn	Leu	Ala	He	Thr	Leu	Arg	Lys	Ala	Gly	Lys
			180					185					190		
Thr	Ala	Asp	Glu	lle	Cys	Lys	Ala	He	His	Val	Phe	Asp	Leu	Lys	Thr
		195					200					205			
Leu	Pro	Val	Asp	Phe	Val	Glu	Cys	Leu	Met	Arg	Phe	Leu	Pro	Thr	Glu
	210					215					220				
Asn	Glu	Val	Lys	Val	Leu	Arg	Leu	Tyr	Glu	Arg	Glu	Arg	Lys	Pro	Leu
225					230					235					240
Glu	Asn	Leu	Ser	Asp	Glu	Asp	Arg	Phe	Met	Met	Gln	Phe	Ser	Lys	He
				245					250					255	
Glu	Arg	Leu	Met	Gln	Lys	Met	Thr	lle	Met	Ala	Phe	lle	Gly	Asn	Phe
			260					265					270		
Ala	Glu	Ser	Ile	Gln	Met	Leu	Thr	Pro	Gln	Leu	His	Ala	lle	Ile	Ala
		275					280					285			
Ala	Ser	Val	Ser	lle	Lys	Ser	Ser	Gln	Lys	Leu	Lys	Lys	lle	Leu	Glu
	290					295					300				
He	He	Leu	Ala	Leu	Gly	Asn	Tyr	Met	Asn	Ser	Ser	Lys	Arg	Gly	Ala
305					310					315					320
Val	Tyr	Gly	Phe	Lys	Leu	Gln	Ser	Leu	Asp	Leu	Leu	Leu	Asp	Thr	Lys
				325					330					335	
Ser	Thr	Asp	Arg	Lys	Gln	Thr	Leu	Leu	His	Tyr	Пe	Ser	Asn	Val	Val
			340					345					350		
Lys	Glu	Lys	Tyr	His	Gln	Val	Ser	Leu	Phe	Tyr	Asn	Glu	Leu	His	Tyr
		355					360					365			
Val	Glu	Lys	Ala	Ala	Ala	Val	Ser	Leu	Glu	Asn	Val	Leu	Leu	Asp	Val
	370					375					380				
Lys	Glu	Leu	Gln	Arg	Gly	Met	Asp	Leu	Thr	Lys	Arg	Glu	Tyr	Thr	Met
385					390					395					400
His	Asp	His	Asn	Thr	Leu	Leu	Lys	Glu	Phe	He	Leu	Asn	Asn		Gly
				405					410					415	
Lys	Leu	Lys		Leu	Gln	Asp	Asp	Ala	Lys	He	Ala	Gln	Asp	Ala	Phe
			420					425					430		

Asp Asp Val Val Lys Tyr Phe Gly Glu Asn Pro Lys Thr Thr Pro Pro 435 440 Ser Val Phe Phe Pro Val Phe Val Arg Phe Val Lys Ala Tyr Lys Gln 450 455 Ala Glu Glu Glu Asn Glu Leu Arg Lys Lys Gln Glu Gln Ala Leu Met 470 475 Glu Lys Leu Glu Gln Glu Ala Leu Met Glu Gln Gln Asp Pro Lys 485 490 Ser Pro Ser His Lys Ser Lys Arg Gln Gln Gln Glu Leu Ile Ala Glu 500 505 Leu Arg Arg Arg Gln Val Lys Asp Asn Arg His Val Tyr Glu Gly Lys 520 Asp Gly Ala Ile Glu Asp Ile Ile Thr Ala Leu Lys Lys Asn Asn Ile 530 535 540 Thr Lys Phe Pro Asn Val His Ser Arg Val Arg Ile Ser Ser Ser Thr 550 555 560 Pro Val Val Glu Asp Thr Gln Ser 565

<210> 4855

<211> 415

<212> PRT

<213> Homo sapiens

<400> 478

Met Thr Leu Ala Ala Ser Ser Gln Arg Ser Gln Ile Ile Arg Ser Lys

1 5 10 15

Phe Arg Ser Val Leu Gln Leu Arg Ile His Arg Arg Asn Gln Glu Gln
20 25 30

lle Ser Asp Pro Asp Pro Trp lle Ser Ala Ser Asp Pro Pro Leu Ala 35 40 45

Pro Ala Leu Pro Ser Gly Thr Ala Pro Phe Leu Phe Ser Pro Gly Val
50 55 60

Leu Leu Pro Glu Pro Glu Tyr Cys Pro Pro Trp Arg Ser Pro Lys Lys
65 70 75 80

G	lu	Ser	Pro	Lys	He	Ser	Gln	Arg	Trp	Arg	Glu	Ser	Lys	Pro		G1 y
					85					90					95	
A	sn	Leu	Thr	Tyr	His	Gln	Tyr	Met	Pro	Pro	Glu	Pro	Arg		Gly	Ser
				100					105					110		
A	rg	Ala	Asp	Pro	Gln	Ala	Glu	Gly	Ser	Ala	Leu	Gly	Pro	Pro	Gly	Pro
			115					120					125			
S	er	Leu	Trp	Glu	Gly	Thr	Asp	Ser	Gln	Gln	Pro	His	Pro	Arg	Met	Lys
		130					135					140				
P	ro	Ser	Pro	Leu	Thr	Pro	Cys	Pro	Pro	Gly	Val	Pro	Ser	Pro	Ser	Pro
1	45					150					155					160
P	ro	Pro	His	Lys	Leu	Glu	Leu	Gln	Thr	Leu	Lys	Leu	Glu	Glu	Leu	Thr
					165					170					175	
V	al	Ser	Glu	Leu	Arg	Gln	Gln	Leu	Arg	Leu	Arg	Gly	Leu	Pro	Val	Ser
				180					185					190		
G	l y	Thr	Lys	Ser	Met	Leu	Leu	Glu	Arg	Met	Arg	Gly	Gly	Ala	Pro	Pro
			195					200					205			
A	rg	Glu	Arg	Pro	Lys	Pro	Arg	Arg	Glu	Asp	Ser	Pro	Ala	Gly	Ala	Pro
		210					215					220				
T	rp	Pro	Arg	Leu	Lys	Pro	Lys	Ala	Leu	Ala	Ala	Ala	Arg	Arg	Gln	Gly
2	25					230					235					240
S	er	Val	Lys	Pro	Ser	Ala	Ala	Ser	His	Arg	Pro	Pro	Leu	Pro	Arg	Ala
					245					250					255	
A	la	Лѕр	Thr	Pro	Gly	Thr	Ala	Pro	Ala	Pro	Thr	Pro	Thr	Pro	Ala	Pro
				260					265					270		
A	la	Λla	Ala	Pro	Ala	Leu	Thr	Pro	Ser	Ser	Gly	Pro	Gly	Ser	Ala	Ala
			275					280					285			
L	eu	Thr	Leu	Glu	Glu	Glu	Leu	Gln	Glu	Ala	lle	Arg	Arg	Ala	Gln	Leu
		290					295					300				
L	eu	Pro	Asn	Arg	Gly	He	Asp	Asp	He	Leu	Glu	Asp	Gln	Val	Glu	Pro
3	05					310					315					320
A	sp	Asp	Pro	Leu	Pro	Pro	Ile	Pro	Leu	Asp	Phe	Pro	Gly	Ser	Phe	Asp
					325					330					335	
V	al	Leu	Ser	Pro	Ser	Pro	Asp	Ser	Glu	G1 y	Leu	Ser	Ser	Val	Phe	Ser
				340					345					350		
C	or.	Sar	Lou	Pro	Sar	Pro	Thr	Acn	Sar	Sor	Sor	Dro	Sor	Pro	Arce	Acr

Pro Thr Asp Ser Leu Asp Trp Leu Glu Ala Leu Ser Gly Gly Pro Pro Leu Gly Ser Gly Pro Pro Pro Pro Ser Ile Phe Ser Ala Asp Leu Ser Asp Ser Ser Ser Ser Arg Leu Trp Asp Leu Leu Glu Asp Pro Trp <210> 4856

<211> 197

<212> PRT

<213> Homo sapiens

<400> 479

Met Ser Cys Tyr Pro Thr Phe Asp Trp Ala Thr Trp Ala Glu Gly Glu Lys Gly Val Trp Lys Ser Pro Pro Ser Ile Gln Thr Gly Asp Gln Ile Gly Val Arg Glu Glu Leu Leu Asn Ala Leu His Ser Ser Leu Ala Arg Pro Ala Ile Lys Lys His Gln His Pro Lys Gly Lys Lys Arg Arg Ser Arg Glu Lys His Gln Glu Ser Thr Thr Asp Pro Gly Ser Pro Lys

Lys Cys Arg Ala Arg Phe Gly Leu Asn Gln Gln Thr Asp Trp Cys Gly

Pro Cys Arg Arg Lys Lys Cys Ile Arg Tyr Leu Pro Gly Glu Gly

Arg Cys Pro Ser Pro Val Pro Ser Asp Asp Ser Ala Leu Gly Cys Pro

Gly Ser Pro Ala Pro Gln Asp Ser Pro Ser Tyr His Leu Leu Pro Arg

Phe Pro Thr Glu Leu Leu Thr Ser Pro Ala Glu Pro Ala Pro Thr Ser Pro Gly Leu Ser Thr Ala Leu Ser Leu Pro Thr Pro Gly Pro Pro Gln

170 165 · 175 Ala Pro Arg Ser Thr Leu Gln Ser Thr Gln Val Gln Gln Gln Glu Ser 180 190 185 Gln Arg Gln Val Ala 195 <210> 4857 <211> 209 <212> PRT <213> Homo sapiens <400> 480 Met Lys Lys Leu Thr Pro Lys Gln Lys Phe Ser Glu Asp Leu Glu Ser 5 10 Tyr Lys Ile Ser Val Val Met Gln Glu Ser Ala Glu Lys Leu Ser Glu 20 25 Lys Leu His Lys Cys Lys Glu Phe Val Asp Ser Cys Arg Leu Thr Phe 35 45 Pro Thr Ser Gly Asp Glu Tyr Ser Arg Gly Phe Leu Gln Asn Leu Asn Leu Ile Gln Asp Gln Asn Ala Gln Thr Arg Trp Lys Gln Gly Arg Tyr 70 75 Asp Glu Asp Gly Lys Pro Phe Asn Gln Arg Ser Leu Leu Leu Gly His 85 90 Glu Arg lle Leu Thr Arg Ala Lys Ser Tyr Glu Cys Ser Glu Cys Gly 100 105 Lys Val Ile Arg Arg Lys Ala Trp Phe Asp Gln His Gln Arg lle His 115 120 125 Phe Leu Glu Asn Pro Phe Glu Cys Lys Val Cys Gly Gln Ala Phe Arg 135 Gln Arg Ser Ala Leu Thr Val His Lys Gln Cys His Leu Gln Asn Lys 150 155 160 Pro Tyr Arg Cys His Asp Cys Gly Lys Cys Phe Arg Gln Leu Ala Phe 165 170 175

Leu Leu Asn Ile Arg Gly Phe Thr Pro Lys Lys Asn Leu Ile Asn Val

Ala Asn Val Lys Lys Arg Leu Val Arg Ile Gln Pro Leu Phe Asp Ile Arg <210> 4858 <211> 259 <212> PRT <213> Homo sapiens <400> 481 Met Glu Cys His Leu Lys Thr His Tyr Lys Met Glu Tyr Lys Cys Arg lle Cys Gln Thr Val Lys Ala Asn Gln Leu Glu Leu Glu Thr Arg Thr Arg Glu His Arg Leu Gly Asn His Tyr Lys Cys Asp Gln Cys Gly Tyr Leu Ser Lys Thr Ala Asn Lys Leu Ile Glu His Val Arg Val His Thr Gly Glu Arg Pro Phe His Cys Asp Gln Cys Ser Tyr Ser Cys Lys Arg Lys Asp Asn Leu Asn Leu His Lys Lys Leu Lys His Ala Pro Arg Gln Thr Phe Ser Cys Glu Glu Cys Leu Phe Lys Thr Thr His Pro Phe Val Phe Ser Arg His Val Lys Lys His Gln Ser Gly Asp Cys Pro Glu Glu Asp Lys Lys Gly Leu Cys Pro Ala Pro Lys Glu Pro Ala Gly Pro Gly Ala Pro Leu Leu Val Val Gly Ser Ser Arg Asn Leu Leu Ser Pro Leu Ser Val Met Ser Ala Ser Gln Ala Leu Gln Thr Val Ala Leu Ser Ala

Ala His Gly Ser Ser Ser Glu Pro Asn Leu Ala Leu Lys Ala Leu Ala Phe Asn Gly Ser Pro Leu Arg Phe Asp Lys Tyr Arg Asn Ser Asp Phe Ala His Leu Ile Pro Leu Thr Met Leu Tyr Pro Lys Asn His Leu Asp Leu Thr Phe His Pro Pro Arg Pro Gln Thr Ala Pro Pro Ser Ile Pro Ser Pro Lys His Pro Phe Leu Ala Tyr Leu Gly Leu Arg Glu Arg Ala

Glu Thr Val

<210> 4859

<211> 160

<212> PRT

<213> Homo sapiens

<400> 482

Met 11e Ser Ala Tyr Cys Asn Leu His Leu Leu Gly Ser Ser Asn Ser Ser Ala Ser Ala Ser Gln Val Ala Gly 11e Thr Gly Ala Cys Gln His Ala Trp Leu lle Trp Gly Tyr Ser Ser Phe Leu Arg Gln Ser Phe Ala Leu Ser Pro Lys Leu Glu Cys Ser Gly Thr Ile Ser Ala Leu Cys Ser Leu Cys Leu Leu Gly Ser Ser Asp Ser Pro Ser Ser Ala Ser Gln Val Ala Gly lle Thr Gly Ala Cys Tyr His Ala Gln Leu Ile Phe Val Phe Leu Val Gln Met Arg Phe His His Leu Gly Gln Ala Gly Leu Glu Leu Leu Thr Ser Ser Asp Leu Pro Ala Ser Ala Ser Gln Ser Ala Gly Ile

Thr Asp Val Asn His His Ala Gln Pro Leu Trp Gly Tyr Ser Cys Pro Tyr Phe Ser Phe Ile Val Leu Tyr Lys Lys Val Val Leu Leu Tyr Leu <210> 4860 <211> 276 <212> PRT <213> Homo sapiens <400> 483 Met Asn Val Gln Ser Ser Arg Ser His Ala Ile Phe Thr Ile His Leu Cys Gln Met Arg Met Cys Thr Gln Pro Asp Leu Val Asn Glu Ala Val Thr Gly Leu Pro Asp Gly Thr Pro Pro Ser Ser Glu Tyr Glu Thr Leu Thr Ala Lys Phe His Phe Val Asp Leu Ala Gly Ser Glu Arg Leu Lys Arg Thr Gly Ala Thr Gly Glu Arg Ala Lys Glu Gly Ile Ser Ile Asn Cys Gly Leu Val Gly Thr Trp Trp Val 11e Ser Pro Ser Thr His Thr Ala Gln Ser Gln Thr Pro Pro Arg Pro Cys Ser Pro Thr Pro Ala Glu Ser Ser Leu Asp Ile Leu Pro Ile Ser Pro Ser Leu Ser Gly Val Thr Pro Val Pro Ser Phe Pro Ile Ser Pro Thr Tyr Pro Cys Ala Tyr Lys

 Pro
 Thr
 G1n
 G1y
 Ala
 Arg
 G1y
 Leu
 Leu
 G1y
 Pro
 Lys
 Arg
 G1n
 Leu
 Ala

 145
 150
 150
 155
 155
 155
 160
 160

 Leu
 G1y
 Asp
 G1n
 Ser
 Lys
 Lys
 Val
 Val

 His
 Val
 Ser
 Tyr
 Arg
 Asp
 Ser
 Lys
 Leu
 Thr
 Arg
 Leu
 G1n
 Asp
 Ser

 185
 180
 185
 185
 185
 190
 190
 190

```
Leu Gly Gly Asn Ser Gln Thr Ile Met Ile Ala Cys Val Ser Pro Ser
        195
                                                 205
Asp Arg Asp Phe Met Glu Thr Leu Asn Thr Leu Lys Tyr Ala Asn Arg
    210
                        215
                                             220
Ala Arg Asn Ile Lys Asn Lys Val Val Val Asn Gln Asp Lys Thr Ser
                    230
                                         235
Gln Gln Ile Ser Ala Leu Arg Ala Glu Ile Ala Arg Leu Gln Met Glu
                245
                                     250
                                                         255
Leu Met Glu Tyr Lys Ala Val Ser Met Leu Leu Gly Ile Ala Gln His
            260
                                 265
                                                     270
Ser Pro Trp Ser
        275
<210> 4861
<211> 159
<212> PRT
<213> Homo sapiens
<400> 484
Met Gln Gly Ala Glu Leu His Pro Arg Cys His Ala Trp Val Ser Val
                                      10
Phe Ala Leu Gly Leu Thr Leu Lys Arg Thr Gly Gln Pro Gln Ala Pro
             20
                                 25
                                                      30
Ser Thr Gly Asp Leu Pro Ser Gly Pro Ala Gly Lys Ala Gly Ser Arg
Glu Leu Arg Gly Glu Pro Pro Thr Trp Val Trp His Pro Pro 11e Ser
     50
                         55
                                              60
Asp Asn Phe His Gln Glu Asp Ser Lys Arg Pro His Val Cys Phe Asp
                     70
                                          75
65
                                                              80
Gly Glu Asp Pro Glu Val Asp Gly Phe Arg Gly Cys Pro Leu Asp Gly
                 85
                                      90
Glu Leu Gly Pro Cys Gly Lys Ser His Ala Ser Phe Gly Arg Pro Arg
            100
                                 105
Pro Val Ala Pro Thr Leu His Pro His Pro Tyr Pro Cys Ser Cys Cys
```

Pro Ser Ile Lys Pro Pro Leu Arg Gly Leu Thr Gly Arg Gly Gln
130

135

140

Ala Trp Arg Gly Ser His Tyr Arg Thr Pro Gly His Met Thr Arg
145

150

155

<210> 4862

<211> 781

<212> PRT

<213> Homo sapiens

<400> 485

Met Lys Ser Ser Gly Pro Val Glu Arg Leu Leu Arg Ala Leu Gly Arg

1 5 10 15

Arg Asp Ser Ser Arg Ala Ala Ser Arg Pro Arg Lys Ala Glu Pro His 20 25 30

Ser Phe Arg Glu Lys Val Phe Arg Lys Lys Pro Pro Val Cys Ala Val 35 40 45

Cys Lys Val Thr Ile Asp Gly Thr Gly Val Ser Cys Arg Val Cys Lys
50 55 60

Val Ala Thr His Arg Lys Cys Glu Ala Lys Val Thr Ser Ala Cys Gln 65 70 75 80

Ala Leu Pro Pro Val Glu Leu Arg Arg Asn Thr Ala Pro Val Arg Arg 85 90 95

lle Glu His Leu Gly Ser Thr Lys Ser Leu Asn His Ser Lys Gln Arg 100 105 110

Ser Thr Leu Pro Arg Ser Phe Ser Leu Asp Pro Leu Met Glu Arg Arg 115 120 125

Trp Asp Leu Asp Leu Thr Tyr Val Thr Glu Arg Ile Leu Ala Ala Ala 130 135 140

Phe Pro Ala Arg Pro Asp Glu Gln Arg His Arg Gly His Leu Arg Glu 145 150 155 160

Leu Ala His Val Leu Gln Ser Lys His Arg Asp Lys Tyr Leu Leu Phe 165 170 175

Asn Leu Ser Glu Lys Arg His Asp Leu Thr Arg Leu Asn Pro Lys Val 180 185 190

Gln	Asp	Phe	Gly	Trp	Pro	Glu	Leu	His	Ala	Pro	Pro	Leu	Asp	Lys	Leu
		195					200					205			
Cys	Ser	Пe	Cys	Lys	Ala	Met	Glu	Thr	Trp	Leu	Ser	Ala	Asp	Pro	G1n
	210					215					220				
His	Val	Val	Val	Leu	Tyr	Cys	Lys	Gly	Asn	Lys	G]y	Lys	Leu	Gly	Val
225					230					235					240
He	Val	Ser	Ala	Tyr	Met	His	Tyr	Ser	Lys	He	Ser	Ala	G1 y	Ala	Asp
				245					250					255	
Gln	Ala	Leu	Ala	Thr	Leu	Thr	Met	Arg	Lys	Phe	Cys	Glu	Asp	Lys	Val
			260					265					270		
Ala	Thr	Glu	Leu	Gln	Pro	Ser	Gln	Arg	Arg	Tyr	Ile	Ser	Tyr	Phe	Ser
		275					280					285			
Gly	Leu	Leu	Ser	GIy	Ser	Пе	Arg	Met	Asn	Ser	Ser	Pro	Leu	Phe	Leu
	290					295					300				
His	Tyr	Val	Leu	He	Pro	Met	Leu	Pro	Ala	Phe	Glu	Pro	Gly	Thr	Gly
305					310					315					320
Phe	Gln	Pro	Phe	Leu	Lys	lle	Tyr	Gln	Ser	Met	Gln	Leu	Val	Tyr	Thr
				325					330					335	
Ser	Gly	Val	Tyr	His	He	Ala	Gly	Pro	Gly	Pro	Gln	Gln	Leu	Cys	Ile
			340					345					350		
Ser	Leu	Glu	Pro	Ala	Leu	Leu	Leu	Lys	Gly	Asp	Val	Met	Va]	Thr	Cys
		355					360					365			
Tyr	His	Lys	Gly	Gly	Arg	Gly	Thr	Asp	Arg	Thr	Leu	Val	Phe	Arg	Val
	370					375					380				
Gln	Phe	His	Thr	Cys	Thr	lle	His	Gly	Pro	Gln	Leu	Thr	Phe	Pro	Lys
385					390					395					400
Asp	Gln	Leu	Asp	Glu	Ala	Trp	Thr	Asp	Glu	Arg	Phe	Pro	Phe	Gln	Ala
				405					410					415	
Ser	Val	Glu	Phe	Va]	Phe	Ser	Ser	Ser	Pro	Glu	Lys	lle	Lys	Gly	Ser
			420					425					430		
Thr	Pro	Arg	Asn	Asp	Pro	Ser	Val	Ser	Val	Asp	Tyr	Asn	Thr	Thr	Glu
		435					440					445			
Pro	Ala	Val	Arg	Trp	Asp	Ser	Tyr	Glu	Asn	Phe	Asn	Gln	His	His	Glu
	450					455					460				
Asp	Ser	Val	Asp	Gly	Ser	Leu	Thr	His	Thr	Arg	Gly	Pro	Leu	Asp	Gly
465					470					475					480

Ser	Pro	Tyr	Ala	Gln	Val	Gln	Arg	Pro	Pro	Arg	Gln	Thr	Pro	Pro	Ala
				485					490					495	
Pro	Ser	Pro	Glu	Pro	Pro	Pro	Pro	Pro	Met	Leu	Ser	Val	Ser	Ser	Asp
			500					505					510		
Ser	Gly	His	Ser	Ser	Thr	Leu	Thr	Thr	Glu	Pro	Ala	Ala	Glu	Ser	Pro
		515					520					525			
Gly	Arg	Pro	Pro	Pro	Thr	Λla	Ala	Glu	Arg	Gln	Glu	Leu	Asp	Arg	Leu
	530					535					540				
Leu	Gly	Gly	Cys	Gly	Val	Ala	Ser	Gly	Gly	Arg	Gly	Ala	Gly	Arg	Glu
545					550					555					560
Thr	Ala	He	Leu	Asp	Asp	Glu	Glu	Gln	Pro	Thr	Val	Gly	G1 y	Gly	Pro
				565					570					575	
His	Leu	Gly	Val	Tyr	Pro	Gly	His	Arg	Pro	Gly	Leu	Ser	Arg	His	Cys
			580					585					590		
Ser	Cys	Arg	Gln	Gly	Tyr	Arg	Glu	Pro	Cys	Gly	Val	Pro	Asn	Gly	Gly
		595					600					605			
Tyr	Tyr	Arg	Pro	Glu	Gly	Thr	Leu	Glu	Arg	Arg	Arg	Leu	Ala	Tyr	Gly
	610					615					620				
Gly	Tyr	Glu	Gly	Ser	Pro	GIn	Gly	Tyr	Ala	Glu	Ala	Ser	Met	Glu	
625					630					635					640
Arg	Arg	Leu	Cys	Arg	Ser	Leu	Ser	Glu	Gly	Leu	Tyr	Pro	Tyr		Pro
				645					650					655	
Glu	Met	Gly		Pro	Ala	Thr	Gly			Gly	Tyr	Arg	Ala	Ser	Ser
			660					665					670		
Ala	Ala			Cys	Ser	Pro			Thr	Pro	Ala		Val	His	Phe
		675					680					685			
Lys			Ala	Gln	Gly			Leu	Thr	Asp			Arg	Lys	Leu
	690					695			C	7.7	700		C	C	T.I
	Phe	Arg	Arg	His			Val	Asn	Ser			Phe	Ser	Ser	
705		0.7			710		TI		D	715		TI.	T1 .	C	720
Asp	Pro	GIn	Asp			Irp	Ihr	Asn			СТУ	ınr	Thr		
1.1	131	C I	ra)	725		,	1 -	D	730		. D	Т	. C1.	735	
ile	Phe	61y			ΑТа	Lys	Lys			ser	rro	ırp	Glu 750		val
C	11.		740		C1	1	Λ	745		C1.	D.	д1	750	_	71 ~
LVC	1110	า คบ	Phe	Ala	1: [11	-L.eu	Asn	rro	- ASD	- GIN	rro) ATA	ı G1 v	_ATA	116

.

755 760 765

Val Thr Phe 11e Thr Lys Val Leu Leu Gly Gln Arg Lys
770 775 780

<210> 4863

<211> 114

<212> PRT

<213> Homo sapiens

<400> 486

Met lle Lys Glu Leu Thr Gln Asn Leu Asn Thr Phe Phe lle Glu Phe 1 5 10 15

lle Tyr Gly Cys Leu Leu Phe Val Glu Phe Phe Ser Ser Leu Phe Ser 20 25 30

Phe Phe Phe Ser Phe Leu Ser Ser Leu Leu Phe Ser Phe Phe Ser Val

Ala Gln Ala Gly Val Gln Trp His Asp Leu Gly Ser Leu Gln Pro Pro 50 55 60

Pro Pro Arg Phe Lys His Leu Ser Cys Leu Ser Leu Pro Ser Ser Trp 65 70 75 80

Asp Tyr Arg Cys Pro Pro Pro Ser Pro Ala Asn Phe Cys 11e Phe Ser 85 90 95

Arg Asp Arg Val Ser Pro Cys Trp Leu Gly Trp Ser Arg Thr Ser Asp 100 105 110

Leu Lys

<210> 4864

<211> 141

<212> PRT

<213> Homo sapiens

<400> 487

Met Pro Val 11e Pro Ala Leu Gly Glu Ala Lys Gly Glu Val Leu Pro

5 10 Pro Gly Asp Thr Thr Thr Ile 11e Pro Leu Asn Trp Met Leu Lys Ser 20 25 30 Pro Pro Gly His Phe Gly Leu Leu Leu Leu Leu Ser Gln Gln Ala Lys 40 Asn Gly Val Met Val Leu Ala Gly Val Thr Asp Pro Glu Tyr Gln Asp 55 60 Glu Ile Ser Leu Leu His Asn Glu Gly His Leu Lys Glu Val Lys 70 75 65 80 Met Glu Gly Ala Arg Leu Gly Leu Pro Gly Arg Ala Glu Ser Leu Glu 85 90 His Gln Val Gln Ser His Leu Asn Met Ile Ala Gln Ser Gln Arg Thr 105 Phe Gln Lys Lys Asp Ala Gly Lys Ala Ile lle Leu Ser, Lys Leu Thr 115 120 125 Gln Glu Gln Lys Thr Lys His Cys Met Phe Ser Leu His 135 140

<210> 4865

<211> 731

<212> PRT

<213> Homo sapiens

<400> 488

Met His Leu Arg Leu Arg Pro Glu Val Gly Arg Ser Arg Ala Arg Ser

1 5 10 15

Gly Glu Pro Ala Gly Ser Ala Ala Ala Arg Glu Val Met Ala Ala Ala 20 25 30

Gly Ser Gly Ser Ser Ala Ser Arg Gly Phe Tyr Phe Asn Thr Val Leu
50 55 60

Ser Leu Ala Arg Ser Leu Ala Val Gln Arg Pro Ala Ser Leu Glu Lys 65 70 75 80

Val Arg Lys Leu Cys Met Cys Pro Val Asp Phe His Gly 11e Phe

				85					90					95	
Gln	Leu	Asp	Glu	Arg	Arg	Arg	Asp	Ala	Val	He	Ala	Leu	Gly	He	Phe
			100					105					110		
Leu	He	Glu	Ser	Asp	Leu	G1n	His	Lys	Asp	Cys	Val	Val	Pro	Tyr	Leu
		115					120					125			
Leu	Arg	Leu	Leu	Lys	Gly	Leu	Pro	Lys	Val	Tyr	Trp	Val	Glu	Glu	Ser
	130					135					140				
Thr	Ala	Arg	Lys	Gly	Arg	Gly	Ala	Leu	Pro	Val	Ala	Glu	Ser	Phe	Ser
145					150					155					160
Phe	Cys	Leu	Val	Thr	Leu	Leu	Ser	Asp	Val	Ala	Tyr	Arg	Asp	Pro	Ser
				165					170					175	
Leu	Arg	Asp	Glu	He	Leu	Glu	Va]	Leu	Leu	Gln	Va]	Leu	His	Val	Leu
			180					185					190		
Leu	Gly	Met	Cys	Gln	Ala	Leu	Glu	He	Gln	Asp	Lys	Glu	Tyr	Leu	Cys
		195					200					205			
Lys	Tyr	Ala	lle	Pro	Cys	Leu	lle	Gly	11e	Ser	Arg	Ala	Phe	Gly	Arg
	210					215					220				
Tyr	Ser	Asn	Met	Glu	Glu	Ser	Leu	Leu	Ser	Lys	Leu	Phe	Pro	Lys	lle
225					230					235					240
Pro	Pro	His	Ser	Leu	Arg	Val	Leu	Glu	Glu	Leu	Glu	Gly	Val	Arg	Arg
				245					250					255	
Arg	Ser	Phe	Asn	Asp	Phe	Arg	Ser	lle	Leu	Pro	Ser	Asn	Leu	Leu	Thr
			260					265					270		
Val	Cys	Gln	Glu	Gly	Thr	Leu	Lys	Arg	Lys	Thr	Ser	Ser	Val	Ser	Ser
		275					280					285			
He	Ser	Gln	Val	Ser	Pro	Glu	Arg	Gly	Met	Pro	Pro	Pro	Ser	Ser	Pro
	290					295					300				
Gly	Gly	Ser	Ala	Phe	His	Tyr	Phe	Glu	Ala	Ser	Cys	Leu	Pro	Asp	Gly
305					310					315					320
Thr	Ala	Leu	Glu		Glu	Tyr	Tyr	Phe		Thr	He	Ser	Ser	Ser	Phe
				325					330					335	
Ser	Va]	Ser		Leu	Phe	Asn	GI y		Thr	Tyr	Lys	Glu		Asn	11e
			340					345					350		
Pro	Leu		Met	Leu	Arg	Glu		Leu	Asn	Leu	Val		Lys	lle	Val
		355			_	_	360					365	_		
Glu	Glu	Ala	Val	Leu	Lys	Ser	Leu	Asp	Ala	He	Val	Ala	Ser	Val	Met

	370					375					380				
Glu	Ala	Asn	Pro	Ser	Ala	Asp	Leu	Tyr	Tyr	Thr	Ser	Phe	Ser	Asp	Pro
385					390					395					400
Leu	Tyr	Leu	Thr	Met	Phe	Lys	Met	Leu	Arg	Asp	Thr	Leu	Tyr	Tyr	Met
				405					410					415	
Lys	Asp	Leu	Pro	Thr	Ser	Phe	Val	Lys	Glu	lle	His	Asp	Phe	Val	Leu
			420					425					430		
Glu	Gln	Phe	Asn	Thr	Ser	Gln	Gly	Glu	Leu	Gln	Lys	He	Leu	His	Asp
		435					440					445			
Ala	Asp	Arg	He	His	Asn	Glu	Leu	Ser	Pro	Leu	Lys	Leu	Arg	Cys	Gln
	450					455					460				
Ala	Asn	Ala	Ala	Cys	Val	Asp	Leu	Met	Val	Trp	Ala	Val	Lys	Asp	Glu
465					470					475					480
Gln	Gly	Ala	Glu	Asn	Leu	Cys	He	Lys	Leu	Ser	Glu	Lys	Leu	Gln	Ser
				485					490					495	
Lys	Thr	Ser	Ser	Lys	Va]	He	Ile	Ala	His	Leu	Pro	Leu	Leu	lle	Cys
			500					505					510		
Cys	Leu	Gln	Gly	Leu	Gly	Arg	Leu	Cys	Glu	Arg	Phe	Pro	Val	Val	Val
		515					520					525			
llis	Ser	Val	Thr	Pro	Ser	Leu	Arg	Asp	Phe	Leu	Val	He	Pro	Ser	Pro
	530					535					540				
Val	Leu	Val	Lys	Leu	Tyr	Lys	Tyr	llis	Ser	Gln	Tyr	llis	Thr	Val	Ala
545					550					555					560
Gly	Asn	Asp	lle		He	Ser	Val	Thr		Glu	llis	Ser	Glu		Thr
				565					570					575	
Leu	Asn	Val		Ser	Gly	Lys	Lys		Gln	Pro	Ser	Met		Glu	Gln
			580					585	_		_		590		
Leu	Arg		lle	Ala	He	Asp		He	Cys	Arg	Cys		Lys	Ala	GIy
		595		_			600					605			
Leu		Val	Asp	Pro	Val		Val	GIu	Ala	Phe	Leu	Ala	Ser	Leu	Ser
	610		æ			615	6.1	c			620			. 1	
	Arg	Leu	lyr	116		GIn	GIU	Ser	Asp		Asp	Ala	ms	Leu	
625 D			T)	7.1	630	4.7	,	C I	112.	635		171	A 1 .	1	640
Pro	лѕр	H1S	ınr		Arg	Ala	Leu	GIŸ		116	Ala	vai	Ala		arg
۸	TL	Dass	1	645	Mat	C1	Dwa	11.	650	C1.5	He	Lau	C1.	655	Lvc
ASI	107	110	1. V S	101	MIG. I	(1111	11 (1)	1 1 1-3	1.61	0110	1 1 4-3	1.1-1.1	11111	11111	1. V.S

Phe Cys Gln Pro Pro Ser Pro Leu Asp Val Leu 11e 11e Asp Gln Leu Gly Cys Leu Val lle Thr Gly Asn Gln Tyr lle Tyr Gln Glu Val Trp Asn Leu Phe Gln Gln Ile Ser Val Lys Ala Ser Ser Val Val Tyr Ser Ala Thr Lys Asp Tyr Lys Asp His Gly Tyr Arg <210> 4866

<211> 288

<212> PRT

<213> Homo sapiens

<400> 489 Met Ser Arg Glu Leu Ala Pro Leu Leu Leu Leu Leu Leu Ser Ile His Ser Ala Leu Ala Met Arg Ile Cys Ser Phe Asn Val Arg Ser Phe Gly Glu Ser Lys Gln Glu Asp Lys Asn Ala Met Asp Val Ile Val Lys Val lle Lys Arg Cys Asp lle Ile Leu Val Met Glu Ile Lys Asp Ser Asn Asn Arg lle Cys Pro Ile Leu Met Glu Lys Leu Asn Arg Asn Ser Arg Arg Gly 11e Thr Tyr Asn Tyr Val 11e Ser Ser Arg Leu Gly Arg Asn Thr Tyr Lys Glu Gln Tyr Ala Phe Leu Tyr Lys Glu Lys Leu Val Ser Val Lys Arg Ser Tyr His Tyr His Asp Tyr Gln Asp Gly Asp Ala Asp

Val Phe Ser Arg Glu Pro Phe Val Val Trp Phe Gln Ser Pro His Thr

Ala Val Lys Asp Phe Val lle lle Pro Leu His Thr Thr Pro Glu Thr

150 155 160 145 Ser Val Lys Glu Ile Asp Glu Leu Val Glu Val Tyr Thr Asp Val Lys 170 175 165 His Arg Trp Lys Ala Glu Asn Phe Ile Phe Met Gly Asp Phe Asn Ala 180 185 Gly Cys Ser Tyr Val Pro Lys Lys Ala Trp Lys Asn Val Arg Leu Arg 200 205 Thr Asp Pro Arg Phe Val Trp Leu Ile Gly Asp Gln Glu Asp Thr Thr 220 210 215 Val Lys Lys Ser Thr Asn Cys Ala Tyr Asp Arg Ile Val Leu Arg Gly 230 235 Gln Glu Ile Val Ser Ser Val Val Pro Lys Ser Asn Ser Val Phe Asp 245 250 255 Phe Gln Lys Ala Tyr Lys Leu Thr Glu Glu Glu Val Arg Leu Pro Ser 260 265 270 Cys Leu Ser Met Pro Leu Ser Trp Lys Asp Glu Leu Ala Trp Ala Thr 275 280 285

<210> 4867

<211> 143

<212> PRT

<213> Homo sapiens

<400> 490

Met Pro Ser Thr Pro Ser Ser Arg Arg Ser Arg Asn Ser Ser Arg Ser 1 5 10 15

Pro Pro Gln Thr Ala Asp Cys Trp Ala Ser Ser Lys Ala Arg Thr Pro 20 25 30

Ser Ser Thr Glu Leu Gln Pro Cys Leu Glu Gly Arg Pro Pro Ser Cys 35 40 45

His Pro Asp Leu Ser Gly Ser Pro Phe Pro Pro Ser Leu Asp Pro Lys
50 55 60

Ser Gly Ala 11e His Gln Glu Cys Arg Pro Leu Val Gly Gly Ala Gly 65 70 75 80

 Cys
 Cys
 Leu
 Pro
 11e
 Gly
 Gln
 Leu
 Leu
 Pro
 Glu
 Met
 Gln
 Arg
 Leu

 Leu
 Glu
 Thr
 Gly
 Trp
 Glu
 Leu
 Ala
 Glu
 Pro
 Arg
 Trp
 Arg
 Gly
 Ser

 Ala
 Pro
 Gly
 His
 Phe
 Thr
 Gly
 Arg
 Glu
 Gly
 Gly
 Lys
 Lys
 Lys
 Lys
 Lys
 Gly
 Cys
 Lys

 Arg
 Leu
 Trp
 Asp
 Thr
 Asp
 Ser
 Gln
 Asp
 His
 11e
 Cys
 Gly
 Cys

 130
 Trp
 Asp
 Thr
 Asp
 Ser
 Gln
 Asp
 His
 11e
 Cys
 Gly
 Cys

<210> 4868

<211> 281

<212> PRT

<213> Homo sapiens

<400> 491

Met Ala Val Asp Ser Leu Leu Arg Ile Leu Lys Thr Leu Arg Leu Asp
1 5 10 15

Phe Ile Gln Glu Leu Glu Val Phe Tyr Trp Cys Arg Asp Phe Leu Val 20 25 30

Leu Ala Glu Pro Asn Leu Ser Ala Ile Gln Leu Gly Arg lle Phe Asn 35 40 45

Leu Arg Ser Leu Lys Leu Phe Tyr Tyr Lys Trp Ala Phe Ser Ser Trp 50 55 60

Val Arg Arg Pro Ser Ser Tyr Phe Phe Ser Gln Leu Thr Met Leu Gly
65 70 75 80

His Leu Arg Lys Leu His Leu Ser His Ser Tyr Leu Val Gly Lys Leu 85 90 95

His Tyr 11e Leu Ser Cys Leu Trp Val Pro Leu His Ser Leu Glu I1e 100 105 110

Cys Asn Cys Lys Leu Leu Asp Thr Asp 11e Thr Tyr Leu Ser Arg Ser
115 120 125

His His Thr Thr Cys Leu Lys Lys Leu Asp Leu Ser Val Asn Asp Leu 130 135 140

Ser Tyr Met Ile Pro Gly Pro Leu Gly Thr Leu Leu Arg Ala Val Ser 145 150 155 160 Gly Thr Leu Gln His Leu Asp Leu Lys His Cys Trp Leu Lys Asp Ala His Leu Ser Ala Leu Leu Pro Ala Leu Cys Arg Cys Ser His Leu Ser Ser Leu Ser Leu Ser Asp Asn Pro 11e Ser Ser Ala Cys Leu Leu Ser Leu Leu Glu His Thr Met Gly Leu Met Glu Leu Lys Gln Val Leu Tyr Pro Ile Pro Val Asp Cys Cys Ile Tyr Leu His Gly Val Cys Arg Gly Pro Val Asn Glu Asp Lys Leu Cys Gln Leu Gln Ala Glu Ile Gln Lys Gln Leu Gln Ala Met Gln Gln Ala Asp Met Gln Trp Ser Pro Ser Thr Val Phe Ala Tyr Ala Ala Gly Ala Val

<210> 4869

<211> 197

<212> PRT

<213> Homo sapiens

<400> 492

Met Val Gln Ala Pro Cys Leu Val Val Leu Ser Phe His Val Val Leu Ser Leu Gln Val His Arg Ser Gln Glu Leu Gly Phe Trp Asn Leu His Leu Asp Phe Arg Arg Cys Met Glu Met Pro Gly Cys Pro Gly Arg Asn Leu Leu Gln Gly Trp Gly Ser His Gly Glu Pro Leu Leu Gly Gln Cys Arg Arg Glu Met Trp Gly Gly Ser Pro Thr Gln Ser Pro Tyr Trp Gly Thr Val Leu Gln Thr Gln Asn Gly Arg Ser Thr Asp Ser Leu His Cys

Val Pro Gly Lys Ala Thr Asp Thr Gln Cys Gln Pro Met Lys Ala Thr Arg Arg Glu Thr Val Pro Cys Lys Ala Thr Gly Ala Glu Leu Leu Lys 115 120 Thr Met Gly Thr Tyr Leu Leu Tyr Gln His Asp Leu Asp Val Arg His 135 Gly Val Lys Gly Glu His Phe Gly Ala Leu Arg Phe Asp Cys Pro Ala 155 145 150 160 Gly Phe Trp Thr Cys Met Gly Pro Val Ala Pro Leu Phe Trp Pro Val 170 165 Ser Pro Ile Trp Asn Gly Tyr Ile Tyr Pro Met Pro Ile Pro Pro Leu 190 185 Tyr Pro Gly Ser Asn 195

<210> 4870

<211> 218

<212> PRT

<213> Homo sapiens

<400> 493

 Met
 Trp
 Leu
 His
 Arg
 Gly
 Pro
 Leu
 Arg
 Pro
 Pro
 Pro
 Gly
 Val
 Arg
 Trp
 Thr

 Pro
 Trp
 Ala
 Phe
 Leu
 Glu
 Ala
 Cys
 Ser
 Trp
 Gly
 Pro
 Ala
 Leu
 Ser
 Leu

 Leu
 Gly
 Ser
 Gly
 His
 Ser
 Leu
 Pro
 Gly
 Thr
 His
 Glu
 Ala
 Ala
 Ala
 Trp

 Leu
 Gly
 Ser
 Gly
 His
 Ser
 Leu
 Pro
 Gly
 Thr
 His
 Glu
 Ala
 Ala
 Ala
 Trp

 Ser
 Arg
 Gly
 Gly
 Gln
 His
 Gly
 Gly
 Gln
 Ser
 Pro
 Thr
 Gln
 Lys
 Cys
 Lys

 Ser
 Arg
 Gly
 Gly
 Gln
 His
 Gly
 Gln
 Ser
 Pro
 Thr
 Gln
 Lys
 Cys
 Lys

Ser Ser Lys Glu Pro Leu Ala Gln Ala Pro Pro His Gln Gly Phe Ala

65 70 75 80

Asp Val Leu Glu Arg Pro Thr Leu Glu Pro Phe Gly Val Leu Ala Pro
85 90 95

Pro Val Pro Ser Ala Leu Val Glu Ala Ala Ala Thr Ser Pro Pro Gln 100 105 110 Gly Ala Pro Arg Gly Ile Leu Trp Asp Arg Cys Pro Gln Ile Gln Val Leu Glu Gly Gln Arg Val Arg Phe Pro Ser Gln Pro Gln His Pro Ser His Leu Ala Pro Arg Gly Gly Cys Gly Trp Arg Pro Asp Ser Arg Pro Leu Leu Pro Thr Pro Ser Gly Leu Ser Ser Phe Phe Pro Leu Asp Ala Gln Cys Trp Pro Trp Arg Thr Val Ser Trp Arg Met Ala Val Gly Glu Ala Val Phe Val Pro Leu Gln His Pro Pro Leu Leu His Gly Ser Pro lle Pro Lys Leu Leu Pro Gly Pro Leu Leu

<210> 4871

<211> 349

<212> PRT

<213> Homo sapiens

<400> 494

Met Ala Met Ala Phe Thr Asp Leu Leu Asp Ala Leu Gly Ser Met Gly Arg Phe Gln Leu Asn His Thr Ala Leu Leu Leu Leu Pro Cys Gly Leu Leu Ala Cys His Asn Phe Leu Gln Asn Phe Thr Ala Ala Val Pro Pro His His Cys Arg Gly Pro Ala Asn His Thr Glu Ala Ser Thr Asn Asp Ser Gly Ala Trp Leu Arg Ala Thr lle Pro Leu Asp Gln Leu Gly Ala Pro Glu Pro Cys Arg Arg Phe Thr Lys Pro Gln Trp Ala Leu Leu Ser Pro Asn Ser Ser Ile Pro Gly Ala Ala Thr Glu Gly Cys Lys Asp Gly

Trp	Val	Tyr	Asn	Arg	Ser	Val	Phe	Pro	Ser	Thr	He	Val	Met	Glu	Trp
		115					120					125			
Asp	Leu	Val	Cys	Glu	Ala	Arg	Thr	Leu	Arg	Asp	Leu	Ala	Gln	Ser	Val
	130					135					140				
Tyr	Met	Ala	Gly	Val	Leu	Val	Gly	Ala	Ala	Val	Phe	Gly	Ser	Leu	Ala
145					150					155					160
Asp	Arg	Leu	Gly	Cys	Lys	Gly	Pro	Leu	Val	Trp	Ser	Tyr	Leu	Gln	Leu
				165					170					175	
Ala	Ala	Ser	Gly	Ala	Ala	Thr	Ala	Tyr	Phe	Ser	Ser	Phe	Ser	Ala	Tyr
			180					185					190		
Cys	Val	Phe	Arg	Phe	Leu	Met	Gly	Met	Thr	Phe	Ser	Gly	lle	lle	Leu
		195					200					205			
Asn	Ser	Val	Ser	Leu	Val	Ala	Ser	Arg	Val	He	Pro	Leu	Ala	Pro	Pro
	210					215					220				
Ala	Trp	Gln	Val	Pro	Val	Ser	Cys	Thr	Glu	Ser	Ala	Glu	Gly	Gly	Cys
225					230					235					240
Asn	Glu	Arg	Glu	Glu	Gly	Gly	Arg	Gly	Lys	Ala	Asp	Gln	Gly	Gly	Asp
				245					250					255	
Glu	Leu	Leu	His	Pro	Lys	Arg	Val	Cys	Lys	Cys	Leu	His	Leu	Gln	Leu
			260					265					270		
Asn	Leu	Gly	Pro	Leu	Pro	Asn	Pro	Gly	His	Pro	Gln	Gly	His	Met	Leu
		275					280					285			
Ser	His	Gly	Asp	Leu	Arg	Gly	Arg	Gln	Arg	Ser	Ser	Gly	Ser	Gly	Glu
	290					295					300				
Gln	Gly	Gly	Ser	Cys	Glu	Pro	Glu	Ser	Arg	Asn	Gly	Arg	Leu	Pro	Ala
305					310					315					320
Val	Ala	Pro	Ala	Gly	Leu	Gly	Leu	Pro	Phe	Va]	His	Leu	Pro	Leu	Ser
				325					330					335	
Ala	Arg	Glu	Lys	Gln	He	Leu	Ser	Ser	Lys	Phe	Gln	Ser			
			340					345							

<211> 421

<212> PRT

<213≻ Homo sapiens

٠	<400)> 49) 5													
	Met	Arg	Tyr	Ser	Leu	Ser	Pro	Asp	Asn	His	Leu	Glu	Asp	Gly	He	Met
	1				5					10					15	
	Asn	Met	Ala	Asn	Phe	Leu	۸rg	Gly	Phe	Glu	Glu	Lys	Gly	He	Lys	Asn
				20					25					30		
	Asp	Arg	Pro	Glu	Asp	Gln	Leu	Ser	Lys	Glu	Lys	Lys	Lys	lle	Leu	Phe
			35					40					45			
	Ser	Phe	Cys	Glu	Val	Cys	Asn	He	Gln	Leu	Asn	Ser	Лlа	Ala	Gln	Ala
		50					55					60				
	Gln	Val	His	Ser	Asn	Gly	Lys	Ser	His	Arg	Lys	Arg	Val	Lys	Gln	Leu
	65					70					75					80
	Ser	Asp	Gly	Gln	Pro	Pro	Pro	Pro	Ala	Gln	Ala	Ser	Pro	Ser	Ser	Asn
					85					90					95	
	Ser	Ser	Thr	Gly	Ser	Thr	Cys	His	Thr	Thr	Thr	Leu	Pro	Ala	Leu	Val
				100					105					110		
	Arg	Thr	Pro	Thr	Leu	Met	Met	Gln	Pro	Ser	Leu	Asp	lle	Lys	Pro	Phe
			115					120					125			
	Met	Ser	Phe	Pro	Val	Asp	Ser	Ser	Ser	Ala	Val	Gly	Leu	Phe	Pro	Asn
		130					135					140				
	Phe	Asn	Thr	Met	Λsp	Pro	Val	Gln	Lys	Ala	Val	lle	Asn	His	Thr	Phe
	145					150					155					160
	G1 y	Val	Ser	He	Pro	Pro	Lys	Lys	Lys	Gln	Va]	lle	Ser	Cys	Asn	Val
					165					170					175	
	Cys	Gln	Leu	Arg	Phe	Asn	Ser	Asp	Ser	Gln	Ala	Glu	Ala	His	Tyr	Lys
				180					185					190		
	Gly	Ser		His	Ala	Lys	Lys	Val	Lys	Ala	Leu	Asp		Thr	Lys	Asn
			195					200					205			
	Lys		Lys	Met	Val	Pro		Lys	Asp	Ser	Ala		Ala	Asn	Pro	Ser
		210					215					220				
		Ser	He	Thr	Pro		Thr	G1 y	Asn	Asn		Asp	Lys	Ser	Glu	
	225					230	_	_		_	235		_		_	240
	Lys	Gly	Lys	Leu		Ala	Ser	Ser	Ser		Gln	Pro	Ser	Ser		Glu
		0.		D)	245				0.	250	m			Б	255	a :
	Ser	Gly	Ser	Phe	Leu	Leu	Lys	Ser	GIy	Thr	Thr	Pro	Leu	Pro	Pro	Gly

			260					265					270		
Ala	Ala	Thr	Ser	Pro	Ser	Lys	Ser	Thr	Asn	G1 y	Ala	Pro	Gly	Thr	Val
		275					280					285			
Val	Glu	Ser	Glu	Glu	Glu	Lys	Ala	Lys	Lys	Leu	Leu	Tyr	Cys	Ser	Leu
	290					295					300				
Cys	Lys	Va]	Ala	Val	Asn	Ser	Leu	Ser	Gln	Leu	Glu	Ala	His	Asn	Thr
305					310					315					320
Gly	Ser	Lys	His	Lys	Thr	Met	Val	Glu	Ala	Arg	Asn	Gly	Ala	Gly	Pro
				325					330					335	
lle	Lys	Ser	Tyr	Pro	Arg	Pro	Gly	Ser	Arg	Leu	Lys	Met	Gln	Asn	G1 y
			340					345					350		
Ser	Lys	Gly	Ser	Gly	Leu	Gln	Asn	Lys	Thr	Phe	His	Cys	Glu	Пе	Cys
		355					360					365			
Asp	Val	His	Val	Asn	Ser	Glu	lle	Gln	Leu	Lys	Gln	His	lle	Ser	Ser
	370					375					380				
Arg	Arg	His	Lys	Asp	Arg	Val	Ala	Gly	Lys	Pro	Leu	Lys	Pro	Lys	Tyr
385					390					395					400
Thr	Pro	Cys	Lys	Val	lle	Gly	Leu	Leu	Pro	Lys	Pro	Leu	Pro	Pro	Ala
				405					410					415	
Asn	Arg	Gln	Leu	Ser											
			420												
)> 48														
	1> 2														
	2> PI														
<213	3> He	omo :	sapi	ens											
< 400	22 40	36													

Met Gly Lys Asn Leu Lys Gly Ile Leu Thr Thr Glu Asn Ile Leu Ser

lle Asp Asn Ser Val Asn Lys Lys Asp Leu Ser lle Cys Gly Ser Ser

Gly Glu Glu Phe Phe Asn Asn Cys Glu Val Leu Gln Cys Gly Phe Ser

Val Pro Arg Glu Asn lle Arg Thr Arg His Lys Ile Cys Pro Cys Asp

	50					55					60				
Lys	Cys	Glu	Lys	Val	Phe	Pro	Ser	lle	Ser	Lys	Leu	Lys	Arg	His	Tyr
65					70					75					80
Leu	lle	His	Thr	Gly	Gln	Arg	Pro	Phe	Gly	Cys	Asn	He	Cys	Gly	Lys
				85					90					95	
Ser	Phe	Arg	Gln	Ser	Ala	His	Leu	Lys	Arg	His	Glu	Gln	Thr	His	Asn
			100					105					110		
Glu	Lys	Ser	Pro	Tyr	Ala	Ser	Leu	Cys	Gln	Val	Glu	Phe	Gly	Asn	Phe
		115					120					125			
Asn	Asn	Leu	Ser	Asn	His	Ser	Gly	Asn	Asn	Val	Asn	Tyr	Asn	Ala	Ser
	130					135					140				
Gln	Gln	Cys	Gln	Ala	Pro	Gly	Val	Gln	Lys	Tyr	Glu	Val	Ser	Glu	Ser
145					150					155					160
Asp	Gln	Met	Ser	Gly	Val	Lys	Ala	Glu	Ser	Gln	Asp	Phe	He	Pro	Gly
				165					170					175	
Ser	Thr	Gly	Gln	Pro	Cys	Leu	Pro	Asn	Val	Leu	Leu	Glu	Ser	Glu	Gln
			180					185					190		
Ser	Asn	Pro	Phe	Cys	Ser	Tyr	Ser	Glu	His	Gln	Glu	Lys	Asn	Asp	Val
		195					200					205			
Phe	Leu	Tyr	Arg	Cys	Ser	Val	Cys	Ala	Lys	Ser	Phe	Arg	Ser	Pro	Ser
	210					215					220				
Lys	Leu	Glu	Arg	His	Tyr	Leu	He	His	Ala	Gly	Gln	Lys	Pro	Phe	G]u
225					230					235					240
Cys	Ser	Val	Cys	Gly	Lys	Thr	Phe	Arg		Ala	Pro	His	Trp		Arg
				245					250					255	
His	Gln	Leu	Thr	His	Phe	Lys	Glu	Arg	Pro	Gln	Gly	Lys	Val	Val	Λla
			260					265					270		
Leu	Asp	Ser	Val	Met											
		275													

<211> 933

<212> PRT

<213> Homo sapiens

<400)> 49	97													
Met	Ala	Trp	Lys	Thr	Leu	Pro	lle	Tyr	Leu	Leu	Leu	Leu	Leu	Ser	Val
l				5					10					15	
Phe	Val	He	Gln	G1n	Val	Ser	Ser	Gln	Glu	Leu	Ser	Cys	Lys	Gly	Arg
			20					25					30		
Cys	Phe	Glu	Ser	Phe	Glu	Arg	Gly	۸rg	Glu	Cys	Asp	Cys	Asp	Ala	Gln
		35					40					. 45			
Cys	Lys	Lys	Tyr	Asp	Lys	Cys	Cys	Pro	Asp	Tyr	Glu	Ser	Phe	Cys	Ala
	50					55					60				
Glu	Val	His	Asn	Pro	Thr	Ser	Pro	Pro	Ser	Ser	Lys	Lys	Ala	Pro	Pro
65					70					75					80
Pro	Ser	Gly	Ala	Ser	Gln	Thr	lle	Lys	Ser	Thr	Thr	Lys	Arg	Ser	Pro
				85					90					95	
Lys	Pro	Pro	Asn	Lys	Lys	Lys	Thr	Lys	Lys	Val	Пе	Glu	Ser	Glu	Glu
			100					105					110		
He	Thr	Glu	Glu	His	Ser	Val	Ser	Glu	Λsn	Gln	Glu	Ser	Ser	Ser	Ser
		115					120					125			
Ser	Ser	Ser	Ser	Ser	Ser	Ser	Ser	Thr	Ile	Arg	Lys	He	Lys	Ser	Ser
	130					135					140				
Lys	Asn	Ser	Ala	Ala	Asn	Arg	Glu	Leu	Gln	Lys	Lys	Leu	Lys	Val	Lys
145					150					155					160
Asp	Asn	Lys	Lys	Asn	Arg	Thr	Lys	Lys	Lys	Pro	Thr	Pro	Lys	Pro	Pro
				165					170					175	
Val	Val	Asp	Glu	Ala	Gly	Ser	Gly	Leu	Λsp	Asn	Gly	Asp	Phe	Lys	Val
			180					185					190		
Thr	Thr	Pro	Asp	Thr	Ser	Thr	Thr	Gln	His	Asn	Lys	Val	Ser	Thr	Ser
		195					200					205			
Pro	Lys	He	Thr	Thr	Ala	Lys	Pro	He	Asn	Pro	Arg	Pro	Ser	Leu	Pro
	210					215					220				
Pro	Asn	Ser	Asp	Thr	Ser	Lys	Glu	Thr	Ser	Leu	Thr	Val	Asn	Lys	G] u
225					230					235					240
Thr	Thr	Val	Glu		Lys	Glu	Thr	Thr		Thr	Asn	Lys	Gln		Ser
				245					250					255	
Thr	Asp	Gly		Glu	Lys	Thr	Thr		Ala	Lys	Glu	Thr		Ser	He
			260					265					270		
Glu	Lys	Thr	Ser	Ala	Lys	Asp	Leu	Ala	Pro	Thr	Ser	Lys	Val	Leu	Ala

		275					280					285			
Lys	Pro	Thr	Pro	Lys	Ala	Glu	Thr	Thr	Thr	Lys	G1 y	Pro	Ala	Leu	Thr
	290					295					300				
Thr	Pro	Lys	Glu	Pro	Thr	Pro	Thr	Thr	Pro	Lys	Glu	Pro	Ala	Ser	Thr
305					310					315					320
Thr	Pro	Lys	Glu	Pro	Thr	Pro	Thr	Thr	lle	Lys	Ser	Ala	Pro	Thr	Thr
				325					330					335	
Pro	Lys	Glu	Pro	Ala	Pro	Thr	Thr	Thr	Lys	Ser	Ala	Pro	Thr	Thr	Pro
			340					345					350		
Lys	Glu	Pro	Ala	Pro	Thr	Thr	Thr	Lys	Glu	Pro	Ala	Pro	Thr	Thr	Pro
		355					360					365			
Lys	Glu	Pro	Ala	Pro	Thr	Thr	Pro	Glu	Thr	Pro	Pro	Pro	Thr	Thr	Ser
	370					375					380				
Glu	Val	Ser	Thr	Pro	Thr	Thr	Thr	Lys	Glu	Pro	Thr	Thr	He	His	Lys
385					390					395					400
Ser	Pro	Asp	Glu	Ser	Thr	Pro	Glu	Leu	Ser	Ala	Glu	Pro	Thr	Pro	Lys
				405					410					415	
Ala	Leu	Glu	Asn	Ser	Pro	Lys	Glu	Pro	G1 y	Val	Pro	Thr	Thr	Lys	Thr
			420					425					430		
Pro	Ala	Ala	Thr	Lys	Pro	Glu	Met	Thr	Thr	Thr	Ala	Lys	Asp	Lys	Thr
		435					440					445			
Thr	Glu	Arg	Asp	Leu	Arg	Thr	Thr	Pro	Glu	Thr	Thr	Thr	Ala	Λla	Pro
	450														
Lys						455					460				
465	Met	Thr	Lys	Glu	Thr		Thr	Thr	Thr	Glu		Thr	Thr	Glu	Ser
	Met	Thr	Lys	Glu	Thr 470		Thr	Thr	Thr	Glu 475		Thr	Thr	Glu	Ser 480
Lys				Glu Thr	470	Ala				475	Lys				480
Lys					470 Thr	Ala			Thr	475	Lys Thr	Thr			480
	lle	Thr	Ala	Thr	470 Thr	Ala Thr	Gln	Va]	Thr 490	475 Ser	Lys	Thr	Thr	Gln 495	480 Asp
	lle	Thr	Ala	Thr 485 Lys	470 Thr	Ala Thr	Gln	Va]	Thr 490	475 Ser	Lys	Thr	Thr	Gln 495 Ala	480 Asp
Thr	lle Thr	Thr	Ala Phe 500	Thr 485 Lys	470 Thr	Ala Thr	GIn Thr	Val Leu 505	Thr 490 Lys	475 Ser Thr	Lys Thr	Thr	Thr Leu 510	Gln 495 Ala	480 Asp Pro
Thr	lle Thr	Thr	Ala Phe 500	Thr 485 Lys	470 Thr 11e	Ala Thr	GIn Thr	Val Leu 505	Thr 490 Lys	475 Ser Thr	Lys Thr	Thr Thr	Thr Leu 510	Gln 495 Ala	480 Asp Pro
Thr	lle Thr Val	Thr Pro Thr 515	Ala Phe 500 Thr	Thr 485 Lys	470 Thr 11e	Ala Thr Thr Lys	GIn Thr Thr 520	Val Leu 505 Ile	Thr 490 Lys Thr	475 Ser Thr	Lys Thr Thr	Thr Thr Glu 525	Thr Leu 510	Gln 495 Ala Met	480 Asp Pro
Thr	lle Thr Val	Thr Pro Thr 515	Ala Phe 500 Thr	Thr 485 Lys Thr	470 Thr 11e	Ala Thr Thr Lys	GIn Thr Thr 520	Val Leu 505 Ile	Thr 490 Lys Thr	475 Ser Thr	Lys Thr Thr	Thr Glu 525 Thr	Thr Leu 510	Gln 495 Ala Met	480 Asp Pro
Thr Lys Lys	Thr Val Pro 530	Thr Pro Thr 515 Glu	Ala Phe 500 Thr	Thr 485 Lys Thr	470 Thr 11e Lys	Thr Thr Lys Lys 535	Gln Thr Thr 520 Pro	Val Leu 505 He Lys	Thr 490 Lys Thr	475 Ser Thr Thr	Lys Thr Thr Ala 540	Thr Thr Glu 525 Thr	Thr Leu 510 11e	Gln 495 Ala Met Ser	480 Asp Pro Asn
Thr Lys Lys	Thr Val Pro 530 Thr	Thr Pro Thr 515 Glu	Ala Phe 500 Thr	Thr 485 Lys Thr	470 Thr 11e Lys	Thr Thr Lys Lys 535 Gln	Gln Thr Thr 520 Pro	Val Leu 505 He Lys	Thr 490 Lys Thr	475 Ser Thr Thr	Thr Thr Ala 540	Thr Thr Glu 525 Thr	Thr Leu 510 11e	Gln 495 Ala Met Ser	480 Asp Pro Asn

				565					570					575	
Thr	Thr	Pro	Thr	Pro	Arg	Lys	Met	Thr	Ser	Thr	Met	Pro	Glu	Leu	Asn
			580					585					590		
Pro	Thr	Ser	Arg	11e	Ala	Glu	Ala	Met	Leu	Gln	Thr	Thr	Thr	Arg	Pro
		595					600					605			
Asn	Ğln	Thr	Pro	Asn	Ser	Lys	Leu	Val	Glu	Val	Asn	Pro	Lys	Ser	Glu
	610					615					620				
Asp	Ala	Gly	Gly	Ala	Glu	Gly	Glu	Thr	Pro	His	Met	Leu	Leu	Arg	Pro
625					630					635					640
His	Val	Phe	Met	Pro	Glu	Val	Thr	Pro	Asp	Met	Asp	Tyr	Leu	Pro	Arg
				645					650					655	
Val	Pro	Asn	Gln	Gly	lle	He	He	Asn	Pro	Met	Leu	Ser	Asp	Glu	Thr
			660					665					670		
Asn	He	Cys	Asn	Gly	Lys	Pro	Val	Asp	G1 y	Leu	Thr	Thr	Leu	Arg	Asn
		675					680					685			
Gly	Thr	Leu	Val	Ala	Phe	Arg	Gly	His	Tyr	Phe	Trp	Met	Leu	Ser	Pro
	690					695					700				
Phe	Ser	Pro	Pro	Ser	Pro	Ala	Arg	Arg	lle	Thr	Glu	Val	Trp	Gly	lle
705					710					715					720
Pro	Ser	Pro	Ile	Asp	Thr	Val	Phe	Thr	Arg	Cys	Asn	Cys	Glu	Gly	Lys
				725					730					735	
Thr	Phe	Phe	Phe	Lys	Asp	Ser	Gln	Tyr	Trp	Arg	Phe	Thr	Asn	Asp	Ile
			740					745					750		
Lys	Asp	Ala	Gly	Tyr	Pro	Lys	Pro	lle	Phe	Lys	Gly	Phe	Gly	Gly	Leu
		755					760					765			
Thr	Gly	Gln	lle	Val	Ala	Ala	Leu	Ser	Thr	Ala	Lys	Tyr	Lys	Asn	Trp
	770					775					780				
Pro	Glu	Ser	Val	Tyr	Phe	Phe	Lys	Arg	Gly	Gly	Ser	Пе	Gln	Gln	Tyr
785					790					795					800
He	Tyr	Lys	Gln	Glu	Pro	Val	Gln	Lys	Cys	Pro	Gly	Arg	Arg	Pro	Ala
				805					810					815	
Leu	Asn	Tyr	Pro	Val	Tyr	Gly	Glu	Thr	Thr	Gln	v_{al}	Arg	Arg	Arg	Arg
			820					825					830		
Phe	Clu	Aro	Ala	ماآ	Clv	Dro	Sor	Gla	Thr	Hie	Thr	Ha	Ara	110	Gln
	010	111 8	MIG	., 10	O1 y	110	561	OTH	1 111	111.5	111.1	116	AI B	116	0111

Tyr Ser Pro Ala Arg Leu Ala Tyr Gln Asp Lys Gly Val Leu His Asn 855 Glu Val Lys Val Ser Ile Leu Trp Arg Gly Leu Pro Asn Val Val Thr 865 870 875 880 Ser Ala lle Ser Leu Pro Asn lle Arg Lys Pro Asp Gly Tyr Asp Tyr 890 885 Tyr Ala Phe Ser Lys Asp Gln Tyr Tyr Asn Ile Asp Val Pro Ser Arg 905 900 910 Thr Ala Arg Ala Ile Thr Thr Arg Ser Gly Gln Thr Leu Ser Lys Val 915 920 925 Trp Tyr Asn Cys Pro 930

<210> 4875

<211> 216

<212> PRT

<213> Homo sapiens

<400> 498

Met Arg Trp Trp His Arg Cys Gly Gln Pro Gly Thr Asp Val Val Leu 10 Thr Gly Lys Asn Leu Pro Arg Pro Thr Leu Ser Thr Pro Ala Glu Ser 20 Ser Gln Pro Gly Gln Gly Ala His Val Gly Pro Asn Ala His Ala Pro 40 45 His Ser Cys Gly Glu Leu Thr Ala Arg Pro Arg Gly Ala Arg Gly Ala 55 60 Gln Arg Pro Arg Ser Pro Leu Leu Arg Arg Ala His Ser Gln Ala Arg 65 70 75 80 Gly Arg Thr Trp Gly Pro Thr Pro Thr Leu Pro Thr Pro Ala Glu Ser

Ser Gln Pro Gly Gln Gly Ala His Val Gly Pro Asn Ala His Ala Pro 100 105 110

90

His Ser Cys Gly Glu Leu Thr Ala Arg Pro Gly Gly Ala Arg Gly Ala 115 120 125 Gln Arg Pro His Ser Pro Leu Leu Arg Arg Ala His Ser Gln Ala Arg 135 Gly Arg Thr Trp Gly Pro Thr Phe Arg Asn Lys Ile Ile Val Tyr Ser 155 145 150 160 Phe Leu Phe Phe Leu Lys Ser Val Phe Leu Leu Arg Thr Pro Asp Val 165 170 Cys Gly Ser Glu Ala Gln Trp Ala Tyr Ser Tyr Leu Thr Trp Gln Thr 185 Trp Lys Trp Lys Leu Leu Cys Ile Thr Arg Ile Ser Arg Ala Ser Ala 205 195 200 Ser Cys Trp Pro Asp Arg Val Pro 215 210

<210> 4876

<211> 904

<212> PRT

<213> Homo sapiens

<400> 499

Met Ala Glu Ala Leu Leu Ala Cys Cys Pro Gly Asp Gln Lys Pro Gly
1 5 10 15

Ile Leu Ala Arg Leu Lys Asp Ile Lys Ala Gln Trp Glu Glu Thr Val 20 25 30

Thr Tyr Met Thr His Cys His Ser Arg Ile Glu Trp Val Trp Leu His
35 40 45

Trp Ser Glu Tyr Leu Leu Ala Arg Asp Glu Phe Tyr Arg Trp Phe Gln
50 55 60

Lys Met Met Val Thr Leu Glu Pro His 11e Glu Leu Gln Leu Gly Leu 65 70 75 80

Lys Glu Lys Gln Trp Gln Leu Ser His Ala Gln Val Leu Leu His Asn 85 90 95

Val Asp Asn Gln Ala Val Leu Leu Asp Arg Leu Leu Glu Glu Ala Ala 100 105 110

Ser Leu Phe Asn Arg Ile Gly Asp Pro Ser Val Asp Glu Asp Ala Gln 115 120 125

Lys	Arg	Met	Lys	Ala	Glu	Tyr	Asp	Ala	Val	Lys	Ala	Lys	Ala	Gln	Lys
	130					135					140				
Arg	Val	Asp	Leu	Leu	Glu	Gln	Val	Ala	Arg	Glu	His	Glu	Glu	Tyr	Gln
145					150					155					160
Лlа	Gly	Val	Asp	Glu	Phe	Gln	Leu	Trp	Leu	Lys	Ala	Val	Val	Glu	Lvs
				165					170					175	
Val	Asn	Gly	Cys	Leu	Gly	Arg	Asn	Cys	Lys	Leu	Pro	He	Thr	Gln	Arg
			180					185					190		
Leu	Ser	Thr	Leu	Gln	Asp	Ile	Ala	Lys	Asp	Phe	Pro	Arg	Gly	Glu	Glu
		195					200					205			
Ser		Glu	Thr	Leu	Glu		Gln	Ser	Ala	Gly	Val	He	Arg	Asn	Thr
	210					215					220				
	Pro	Leu	Gly	Ala		Lys	He	Thr	G1 y		Leu	Glu	Glu	Met	
225					230				_	235					240
Lys	Val	Leu	Glu		Leu	Arg	Ala	Leu		Glu	Glu	Glu	Glu		Arg
		61	,	245		C		C1	250	T.	C1	61	C1	255	,
Leu	Arg	Gly		Leu	Arg	Ser	Arg		Ala	Trp	Glu	GIn		He	Lys
C1		C1	260	C1.	1	C	C1	265	۸	М	V - 1	1	270	A	1
GIN	Leu		АТА	GIU	Leu	Ser	Glu	Pne	Arg	мет	vai	285	GIN	Arg	Leu
A10	Cln	275	C1v	Lou	Cln	Dro	280 Ala	Λlο	Lvc	Alo	Cly		Clu	Acn	Clu
піа	290	Ulu	Gly	Leu	GIII	295	МІА	пта	Lys	ма	300	1111	Gju	лър	Oju
Len		Ala	His	Trn	Aro		Tyr	Ser	Ala	Thr		Ala	Ala	Len	Ala
305	, 41	1110	1113	116	310	m g	.,1	001	A. G	315	,,, 8	7110	11,10	Dea	320
	Glu	Glu	Pro	Arg		Asp	Arg	Leu	Gln		Gln	Leu	Lvs	Glu	
				325			8		330					335	
lle	Val	Phe	Pro		Asn	Leu	Lys	Pro		Ser	Asp	Ser	Val		Ala
			340				-	345					350		
Thr	He	Gln	Glu	Tyr	Gln	Ser	Leu	Lys	Val	Lys	Ser	Ala	Arg	Leu	Arg
		355					360					365			
Asn	Ala	Ala	Ala	Val	Glu	Leu	Trp	Gln	His	Phe	Gln	Arg	Pro	Leu	G] n
	370					375					380				
Asp	Leu	Gln	Leu	Trp	Lys	Ala	Leu	Ala	Gln	Arg	Leu	Leu	Glu	Val	Thr
385					390					395					400
Ala	Ser	Leu	Pro	Asp	Leu	Pro	Ser	Leu	His	Thr	Phe	Leu	Pro	Gln	Πe
				405					410					415	

Glu	Ala	Ala	Leu	Met	Glu	Ser	Ser	Arg	Leu	Lys	Glu	Leu	Leu	Thr	Met
			420					425					430		
Leu	Gln	Leu	Lys	Lys	Asp	Leu		lle	Gly	He	Phe	Gly	Gln	Glu	Arg
		435					440					445			
Ala		Ala	Leu	Leu	Glu	Gln	Val	Ala	Gly	Ser	Met	Arg	Asp	Arg	Asp
	450					455					460				
Leu	Leu	His	Asn	Ser	Leu	Leu	Gln	Arg	Lys	Ser	Lys	Leu	Gln	Ser	Leu
465					470					475					480
Leu	Ala	Gln	His		Asp	Phe	Gly	Ala	Ala	Phe	Glu	Pro	Leu		Arg
				485					490					495	
Lys	Leu	Leu		Leu	Gln	Val	Arg		G1n	Ala	Glu	Lys		Leu	Gln
			500					505					510		
Arg	Asp		Pro	Gly	Lys	Gln		Gln	Leu	Ser	Arg		Gln	Gly	Leu
		515					520					525			
GIn		Glu	Gly	Leu	Asp		Gly	Ala	G1n	Met		Ala	Ala	Arg	Pro
	530		6.3			535		0.1			540		0.1		
	Val	GIn	Glu	Asn		Asn	His	GIn	His		Met	Asp	GIn	Leu	
545		151	0.1		550	0.1				555			., .		560
Ser	Asp	Phe	GIn		Leu	GIn	Arg	Ser	Leu	Glu	Asp	Leu	Val		Arg
6	Α.	C1	C	565	C1	01	11.	C	570	D)	C	11.2	C1	575	
Çys	Arg	GIn		vai	GIn	Glu	HIS		Thr	Phe	Ser	HIS		Leu	Leu
C1		Δ	580	т	71.	V = 1	<i>V</i> = 1	585	M - 4	C1	1	f	590	A T	112 -
61u	Leu		GIN	ırp	116	vai		ınr	Met	GIN	Lys		Glu	мта	HIS
A 20.07	C1	595	Alo	C1	Dana	Clu	600	110	C1	Con	Cln	605	A10	C1	Dho
AJ		uro	мта	GIY	110	615	ASP	Ala	G] u	261	620	GIU	ма	Glu	rne
Clu	610	Lou	Val	11a	Clu		Dro	Clu	Lys	C1		Cln	Lou	Son	Lou
Gju	MIG	Leu	vai	Ма	Olu	rne	110	uto	LyS	GIU	MIA	0111	Leu	361	Leu
625					630					635					640
	Glu	Ala	Gln	Glv		Leu	Val	Met	Glu		Ser	Ser	Pro	Glu	
• • • •	010	7110	0117	645	ııρ	Lou		MC C	650	15,5	001	001	110	655	OI,
Ala	Ala	Val	Val		Glu	Glu	Len	Arg	Glu	Len	Ala	Glu	Ser		Arg
7170	1110	, (11	660	0111	0,10	014	Lea	665	010	Lea	71.10	ora	670	1. p	6
Ala	Leu	Arø		Leu	Glu	Glu	Ser		Leu	Ser	Len	Tle		Asn	Trn
		675			-10	-14	680			~ ~ .	.,,,,	685	6		12
His	l.eu		Arø	Met	Glu	Val		Ser	Gly	l.v <	Lvs		Val	Phe	Thr
		~ 111	8		. I U				~		-, 0				

	690					695					700				
Asn	Asn	He	Pro	Lys	Ser	Gly	Phe	Leu	He	Asn	Pro	Met	Asp	Pro	Ile
705					710					715					720
Pro	Arg	His	Arg	Arg	Arg	Glu	Glu	Glu	Gly	Ser	His	Glu	Asp	Phe	Ser
				725					730					735	
Gln	Leu	Leu	Arg	Asn	Phe	G1 y	G1n	Trp	Leu	Gln	Val	Glu	Asn	Ser	Lys
			740					745					750		
Leu	Val	Arg	lle	lle	Ala	Met	Arg	Thr	Ser	Thr	Ala	Glu	Asp	Leu	Arg
		755					760					765			
Thr	Arg	Lys	Ser	Lys	Leu	Gln	Glu	Leu	Glu	Ala	Arg	Val	Pro	Glu	G1 y
	770					775					780				
Gln	His	Leu	Phe	Glu	Asn	Leu	Leu	Arg	Leu	Gly	Pro	Ala	Arg	G] y	Thr
785					790					795					800
Ser	Лsp	Glu	Leu	Glu	Asp	Leu	Arg	Tyr	Gln	Trp	Met	Leu	Tyr	Lys	Ser
				805					810					815	
Lys	Leu	Lys	Asp	Ser	Gly	His	Leu	Leu	Thr	Gln	Ser	Ser	Pro	Gly	Glu
			820					825					830		
Pro	Thr	Gly	Phe	Gln	Lys	Thr	Arg	Arg	Trp	Arg	Gly	Leu	Gly	Ser	Leu
		835					840					845			
Phe	Arg	Arg	Ala	Cys	Cys	Val	Ala	Leu	Pro	Leu	Gln	Leu	Leu	Leu	Leu
	850					855					860				
Leu	Phe	Leu	Leu	Leu	Leu	Phe	Leu	Leu	Pro	He	Arg	Glu	Glu	Asp	Arg
865					870					875					880
Ser	Cys	Thr	Leu	Ala	Asn	Asn	Phe	Ala	Arg	Ser	Phe	Thr	Leu	Met	Leu
				885					890					895	
Arg	Tyr	Asn	Gly	Pro	Pro	Pro	Thr								
			900												

<211> 164

<212> PRT

<213> Homo sapiens

<400> 500

Met Ser Glu Pro Ser Pro Cys Pro Pro Glu Gly Val Gln Leu Cys Gly

5 10 15 Pro Ala Leu Tyr Arg Leu Val Pro Cys Val Cys Ala Val Ser Val Cys 25 Leu Leu Pro Gln Val Leu Lys Lys Tyr Pro Arg Leu Trp Cys Met Thr 40 Lys Pro Pro Ser Arg Arg Pro Lys Leu Tyr 11e Val Asn Leu Gln Trp 55 Thr Pro Lys Asp Asp Trp Ala Ala Leu Lys Leu His Gly Lys Cys Asp 70 75 Asp Val Met Arg Leu Leu Met Ala Glu Leu Gly Leu Glu Ile Pro Ala 85 90 Tyr Ser Arg Trp Gln Asp Pro 11e Phe Ser Leu Ala Thr Pro Leu Arg 100 105 Ala Gly Glu Glu Gly Ser His Ser Arg Lys Ser Leu Cys Arg Ser Arg 115 120 125 Glu Glu Ala Pro Pro Gly Asp Arg Gly Ala Pro Leu Ser Ser Ala Pro 135 140 Ile Leu Gly Gly Trp Phe Gly Arg Gly Cys Thr Lys Arg Thr Lys Arg 150 155 Lys Lys Val Thr

<210> 4878

<211> 477

<212> PRT

<213> Homo sapiens

<400> 501

 Met Lys Leu Ile Arg Ser Ser Ser Cys His Val Gly Ser Ala Arg Lys

 1
 5
 10
 15

 Leu His His Trp Arg Ala Gly Gln Thr Glu Pro Gly Trp Ala Gly Thr
 20
 25
 30

 Arg His Asp Ser Pro Val Pro Leu Pro Pro Arg Gly Gly Leu Ala Ala
 35
 40
 45

 Asp Asn Val Leu Leu Ser Ser Asp Gly Ser His Ala Ala Leu Cys Asp

	50					55					60				
Phe	Gly	llis	Ala	Val	Cys	Leu	Gln	Pro	Asp	Gly	Leu	Gly	Lys	Ser	Leu
65					70					75					80
Leu	Thr	G]y	Asp	Tyr	He	Pro	Gly	Thr	Glu	Thr	His	Met	Ala	Pro	Glu
				85					90					95	
Val	Va]	Leu	Gly	Arg	Ser	Cys	Asp	Ala	Lys	Val	Asp	Val	Trp	Se.r	Ser
			100					105					110		
Cys	Cys	Met	Met	Leu	His	Met	Leu	Asn	Gly	Cys	His	Pro	Trp	Thr	Gln
		115					120					125			
Phe	Phe	Arg	Gly	Pro	Leu	Cys	Leu	Lys	He	Ala	Ser	Glu	Pro	Pro	Pro
	130					135					140				
Val	Arg	Glu	He	Pro	Pro	Ser	Cys	Ala	Pro	Leu	Thr	Ala	Gln	Ala	Ile
145					150					155	٠				160
Gln	Glu	Gly	Leu		Lys	Glu	Pro	He	His	Arg	Val	Ser	Ala	Ala	Glu
				165					170					175	
Leu	Gly	Gly		Val	Asn	Arg	Ala		Gln	Gln	Val	G1 y	Gly	Leu	Lys
			180					185					190		
Ser	Pro		Arg	Gly	Glu	Tyr		Glu	Pro	Arg	His		Pro	Pro	Asn
0.1		195			~ `		200					205	0.1		
GIn		Asn	Tyr	His	GIn		Leu	His	Ala	Gln		Arg	Glu	Leu	Ser
D	210		D	61	D	215	D	. 1	61	C.I	220	TI	C1		4.1
	Arg	Ala	Pro	GIÿ		Arg	Pro	Ala	Glu		Ihr	Ihr	Gly	Arg	
225 Dec	1	1	C1	D	230 Page 1	I	D	D	C1	235	D	C1	Dava	Λ	240
Pro	Lys	Leu	GIII		rro	Leu	rro	rro		rro	Pro	Glu	Pro		Lys
Sor	Dro	Pro	Lou	245	Lou	Sor	Lve	Clu	250	Sor	c1v	Mot	Trn	255	Pro
361	110	110	260	1111	Leu	261	Lys	265	Olu	261	Oly	Mer	Trp 270	Giu	110
Leu	Pro	Leu		Ser	Leu	Glu	Pro		Pro	Ala	Arg	Asn	Pro	Ser	Ser
1,50.0	110	275		501	1,cu	O. u	280	,11 G	110	7110	.11 8	285	110	561	001
Pro	Glu		Lvs	Ala	Thr	Val		Glu	Gln	Glu	Leu		Gln	l.eu	Glu
	290	0				295					300				
He		Leu	Phe	Leu	Asn		Leu	Ser	Gln	Pro		Ser	Leu	Glu	Glu
305					310					315					320
	Glu	Gln	He	Leu		Cys	Leu	Ser	He		Ser	Leu	Ser	Leu	
				325		•			330	-				335	
Asn	Asn	Ser	Glu	Lve	Aen	Pro	Sor	Lve	Ala	Sor	Gln	Sor	Ser	Ara	Asn

			340					345					350		
Thr	Leu	Ser	Ser	Gly	Val	His	Ser	Trp	Ser	Ser	Gln	Ala	Glu	Ala	Arg
		355					360					365			
Ser	Ser	Ser	Trp	Asn	Met	Val	Leu	Ala	Arg	Gly	Arg	Pro	Thr	Asp	Thr
	370					375					380				
Pro	Ser	Tyr	Phe	Asn	Gly	Va]	Lys	Val	Gln	He	GIn	Ser	Leu	Asn	Gly
385					390					395					400
Glu	His	Leu	His	lle	Arg	Glu	Phe	llis	Arg	Val	Lys	Val	G1 y	Asp	11e
				405					410					415	
Лlа	Thr	Gly	lle	Ser	Ser	Gln	Пе	Pro	Ala	Ala	Ala	Phe	Ser	Leu	Val
			420					425					430		
Thr	Lys	Asp	Gly	Gln	Pro	Val	Arg	Tyr	Asp	Met	Glu	Va1	Pro	Asp	Ser
		435					440					445			
Gly	He	Asp	Leu	Gln	Cys	Thr	Leu	Ala	Pro	Asp	Gly	Ser	Phe	Ala	Trp
	450					455					460				
Ser	Trp	Arg	Val	Lys	His	Gly	Gln	Leu	Glu	Asn	Arg	Pro			
465					470					475					

<211> 486

<212> PRT

<213> Homo sapiens

<400> 502

 Met
 Gly
 Ser
 Glu
 Lys
 Asp
 Ser
 Glu
 Ser
 Pro
 Arg
 Ser
 Thr
 Ser
 Leu
 His

 1
 5
 5
 6
 10
 10
 10
 15
 15

 Ala
 Ala
 Ala
 Pro
 Asp
 Pro
 Lys
 Cys
 Arg
 Ser
 Gly
 Gly
 Arg
 Arg
 Arg
 Arg

 Leu
 Leu
 Ser
 Cys
 Pro
 His
 Arg
 Ser
 Cys
 Arg
 Arg
 Cys
 Leu
 Arg
 His
 Tyr

 Leu
 Arg
 Leu
 Glu
 Bro
 Glu
 Ser
 Arg
 Val
 Pro
 He
 Ser
 Cys
 Pro
 Glu

 Cys
 Ser
 Glu
 Arg
 Leu
 Arg
 He
 Arg
 Leu
 Leu
 Arg
 Arg

 Leu
 Arg
 Leu
 Arg
 He
 Arg
 Leu
 Leu
 Leu
 Arg
 Arg

 Leu
 Bro
 His
 Arg
 His
 <t

				85					90					95	
Λla	Ser	Asp	Pro	Asp	Cys	Arg	Trp	Cys	Pro	Ala	Pro	Asp	Cys	Gly	Tyr
			100					105					110		
Ala	Val	lle	Ala	Tyr	Gly	Cys	Ala	Ser	Cys	Pro	Lys	Leu	Thr	Cys	Glu
		115					120					125			
Arg	Glu	Gly	Cys	Gln	Thr	Glu	Phe	Cys	Tyr	His	Cys	Lys	Gln	lle	Trp
	130					135					140				
His	Pro	Asn	Gln	Thr	Cys	Asp	Met	Ala	Arg	Gln	GIn	Arg	Ala	Gln	Thr
145					150					155					160
Leu	Arg	Val	Arg	Thr	Lys	His	Thr	Ser	Gly	Leu	Ser	Tyr	Gly	Gln	G1u
				165					170					175	
Ser	Gly	Pro	Asp	Asp	lle	Lys	Pro	Cys	Pro	Arg	Cys	Ser	Ala	Tyr	He
			180					185					190		
lle	Lys	Met	Asn	Asp	Gly	Ser	Cys	Asn	His	Met	Thr	Cys	Ala	Val	Cys
		195					200					205			
Gly	Cys	Glu	Phe	Cys	Trp	Leu	Cys	Met	Lys	Glu	He	Ser	Asp	Leu	His
	210					215					220				
Tyr	Leu	Ser	Pro	Ser	Gly	Cys	Thr	Phe	Trp	Gly	Lys	Lys	Pro	Trp	Ser
225					230					235					240
Arg	Lys	Lys	Lys	He	Leu	Trp	Gln	Leu	Gly	Thr	Leu	He	Gly	Ala	Pro
				245					250					255	
Val	Gly	lle	Ser	Leu	11e	Ala	Gly	He	Ala	Пе	Pro	Ala	Met	Val	lle
			260					265					270		
Gly	He	Pro	Val	Tyr	Val	Gly	Arg	Lys	Пe	llis	Ser	Arg	Tyr	Glu	Gly
		275					280					285			
Arg	Lys	Thr	Ser	Lys	His	Lys	Arg	Asn	Leu	Ala	He	Thr	Gly	Gly	Val
	290					295					300				
	Leu	Ser	Val	He		Ser	Pro	Va]	He		Ala	Val	Ser	Val	
305					310					315				_	320
lle	Gly	Val	Pro		Met	Leu	Ala	Tyr		Tyr	Gly	Val	Val		He
				325					330					335	
Ser	Leu	Cys		G1 y	Gly	Gly	Cys		Val	Ser	Thr	Ala		Gly	Lys
a			340		•			345					350		
Gly	Val		He	Glu	Phe	Asp		Asp	Asp	Gly	Pro		Thr	Val	Ala
		355					360	1 5	6		٥.	365	C	C	
Acn	Λla	Tres	Δνσ	Ala	Lau	Lvc	Acn	Pro	Sor	Lla	Gilve	Glu	Sar	Sor	HIG

Glu Gly Leu Thr Ser Val Leu Ser Thr Ser Gly Ser Pro Thr Asp Gly Leu Ser Val Met Gln Gly Pro Tyr Ser Glu Thr Ala Ser Phe Ala Ala Leu Ser Gly Gly Thr Leu Ser Gly Gly 11e Leu Ser Ser Gly Lys Gly Lys Tyr Ser Arg Leu Glu Val Gln Ala Asp Val Gln Lys Glu 11e Phe Pro Lys Asp Thr Ala Ser Leu Gly Ala Ile Ser Asp Asn Ala Ser Thr Arg Ala Met Ala Gly Ser Ile Ile Ser Ser Tyr Asn Pro Gln Asp Arg Phe Ser Met Ile His Ala

<210> 4880

<211> 492

<212> PRT

<213> Homo sapiens

<400> 503

Met Ala Met Ala Leu Pro Met Pro Gly Pro Gln Glu Ala Val Val Phe Glu Asp Val Ala Val Tyr Phe Thr Arg lle Glu Trp Ser Cys Leu Ala Pro Asp Gln Gln Ala Leu Tyr Arg Asp Val Met Leu Glu Asn Tyr Gly Asn Leu Ala Ser Leu Gly Phe Leu Val Ala Lys Pro Ala Leu Ile Ser Leu Leu Glu Gln Gly Glu Glu Pro Gly Ala Leu Ile Leu Gln Val Ala Glu Gln Ser Val Ala Lys Ala Ser Leu Cys Thr Asp Ser Arg Met Glu

Ala Gly 11e Met Glu Ser Pro Leu Gln Arg Lys Leu Ser Arg Gln Ala

			100					105					110		
Gly	Leu	Pro	Gly	Thr	Val	Trp	Gly	Cys	Leu	Pro	Trp	Gly	His	Pro	Val
		115					120					125			
Gly	Gly	His	Pro	Ala	Pro	Pro	His	Pro	His	Gly	Gly	Pro	Glu	Asp	Gly
	130					135					140				
Ser	Asp	Lys	Pro	Thr	His	Pro	Arg	Ala	Arg	Glu	His	Ser	Ala	Ser	Pro
145					150					155					160
Arg	Val	Leu	Gln	Glu	Asp	Leu	Gly	Arg	Pro	Val	Gly	Ser	Ser	Ala	Pro
				165					170					175	
Arg	Tyr	Arg	Cys	Val	Cys	Gly	Lys	Ala	Phe	Arg	Tyr	Asn	Ser	Leu	Leu
			180					185					190		
Leu	Arg	His	Gln	He	Val	His	Thr	Gly	Ala	Lys	Pro	Phe	Gln	Cys	Thr
		195					200					205			
Glu	Cys	Gly	Lys	Ala	Phe	Lys	Gln	Ser	Ser	lle	Leu	Leu	Arg	His	Gln
	210					215					220				
Leu	Ile	His	Thr	Glu	Glu	Lys	Pro	Phe	Gln	Cys	Gly	Glu	Cys	G] y	Lys
225					230					235					240
Ala	Phe	Arg	Gln	Ser	Thr	Gln	Leu	Ala	Ala	His	His	Arg	Val	His	Thr
				245					250					255	
Arg	Glu	Arg	Pro	Tyr	Ala	Cys	Gly	Glu	Cys	Gly	Lys	Ala	Phe	Ser	Arg
			260					265					270		
Ser	Ser	Arg	Leu	Leu	Gln	His	Gln	Lys	Phe	llis	Thr	Gly	Glu	Lys	Pro
		275					280					285			
Phe	Ala	Cys	Thr	Glu	Cys	Gly	Lys	Ala	Phe	Cys	Arg	Arg	Phe	Thr	Leu
	290					295					300				
Asn	Glu	His	Gly	Arg	He	His	Ser	Gly	Glu	Arg	Pro	Tyr	Arg	Cys	Leu
305					310					315					320
Arg	Cys	Gly	Gln	Arg	Phe	lle	Arg	Gly	Ser	Ser	Leu	Leu	Lys	llis	His
				325					330					335	
Arg	Leu	His	Ala	Gln	Glu	Gly	Ala	Gln	Asp	G1 y	Gly	Ala	Gly	Gln	G] y
			340					345					350		
Ala	Leu	Leu	Gly	Ala	Ala	Gln	Arg	Pro	G1n	Ala	Gly	Asp	Pro	Pro	His
		355					360					365			
Glu	Cys	Pro	Val	Cys	Gly	Arg	Pro	Phe	Arg	His	Asn	Ser	Leu	Leu	Leu
	370					375					380				
Leu	His	Leu	Arg	Leu	His	Thr	Gly	Glu	Lys	Pro	Phe	Glu	Cys	Ala	Glu

390 395 400 385 Cys Gly Lys Ala Phe Gly Arg Lys Ser Asn Leu Thr Leu His Gln Lys 405 410 Ile His Thr Lys Glu Lys Pro Phe Ala Cys Thr Glu Cys Gly Lys Ala 420 425 430 Phe Arg Arg Ser Tyr Thr Leu Asn Glu His Tyr Arg Leu His Ser Gly 440 445 Glu Arg Pro Tyr Arg Cys Arg Ala Cys Gly Arg Ala Cys Ser Arg Leu 450 455 460 Ser Thr Leu Ile Gln His Gln Lys Val His Gly Arg Glu Pro Gly Glu 470 475 480 Asp Thr Glu Gly Arg Arg Ala Pro Cys Trp Ala Ser 485 490

<210> 4881

<211> 232

<212> PRT

<213> Homo sapiens

<400> 504

Met 11e Arg Lys Val Lys Val Glu Asp Glu Asp Gln Glu Ala Glu Glu 1 5 10 15

Glu Val Glu Trp Pro Gln His Leu Ser Leu Leu Pro Ser Pro Phe Pro 20 25 30

Ala Pro Asp Leu Gly His Leu Ala Ala Ala Tyr Lys Leu Glu Pro Gly
35 40 45

Ala Pro Gly Ala Leu Ser Gly Leu Ala Leu Ser Gly Trp Gly Pro Met 50 55 60

Pro Glu Lys Pro Tyr Gly Cys Gly Glu Cys Glu Arg Arg Phe Arg Asp
65 70 75 80

Gln Leu Thr Leu Arg Leu His Gln Arg Leu His Arg Gly Glu Gly Pro 85 90 95

Cys Ala Cys Pro Ser Leu Val Arg Ser Pro Arg Gly Gly Ala Ala Pro 100 105 110

Ala	Leu	Leu	Leu	Ser	Leu	Val	Leu	Thr	Arg	Thr	Leu	Ser	Cys	Pro	Gln
		115					120					125			
Phe	Arg	Glu	Ala	Arg	Ala	Met	Arg	Pro	Pro	Gly	Val	Ser	Lys	Λla	Thr
	130					135					140				
Trp	Ala	Ala	Ala	Arg	Arg	Phe	Gly	Arg	Arg	Gly	Thr	Pro	Val	Ser	Phe
145					150					155					160
Pro	Gln	Cys	Leu	Arg	Pro	His	Ser	Ile	Pro	Ser	Ser	Asp	Leu	Leu	Gly
				165					170					175	
Gln	Arg	Leu	Ser	Glu	Pro	Leu	Leu	Gly	Thr	Ala	Glu	Leu	Lys	Phe	Leu
			180					185					190		
Glu	Gly	Ser	His	Pro	Gly	Ala	Pro	Leu	Glu	Ser	Arg	Tyr	Phe	Pro	Asp
		195					200					205			
Pro	Ala	Arg	Pro	Gln	Pro	Gly	Gln	Glu	Arg	Val	Val	He	Tyr	Val	Leu
	210					215					220				
Lys	Val	Ser	Leu	Lys	Leu	Lys	Ser								
225					230										

<211> 158

<212> PRT

<213> Homo sapiens

<400> 505

Met Gly Arg Ser Pro Arg Lys 11e Asp Gln Phe Cys Asn Ser Ser Asn 1 5 10 15 Met Val His Gly Ser Val Thr Phe Arg Asp Val Ala Ile Asp Phe Ser 25 Gln Glu Glu Trp Glu Cys Leu Gln Pro Asp Gln Arg Thr Leu Tyr Arg 35 40 45 Asp Val Met Leu Glu Asn Tyr Ser His Leu Ile Ser Leu Gly Ser Ser 55 lle Ser Lys Pro Asp Val lle Thr Leu Leu Glu Gln Glu Lys Glu Pro 75 Trp Met Val Val Arg Lys Glu Thr Ser Arg Arg Tyr Pro Asp Leu Glu

90

95

85

<210> 4883

<211> 296

<212> PRT

<213> Homo sapiens

<400> 506

Met Ala Gln Arg Val Lys Phe Pro Thr Asp Thr Leu Gln Glu Leu Leu

1 5 10 15

Asp Val His Ala Ala Cys Glu Arg Glu Ala Ile Ala Ile Phe Met Glu 20 25 30

His Ser Phe Lys Asp Glu Asn Gln Glu Phe Gln Lys Lys Phe Met Glu 35 40 45

Thr Thr Met Asn Lys Lys Gly Asp Phe Leu Leu Gln Asn Glu Glu Ser 50 55 60

Ser Val Gln Tyr Cys Gln Ala Lys Leu Asn Glu Leu Ser Lys Gly Leu 65 70 75 80

Met Glu Ser lle Ser Ala Gly Ser Phe Ser Val Pro Gly Gly His Lys 85 90 95

Leu Tyr Met Glu Thr Lys Glu Arg Ile Glu Gln Asp Tyr Trp Gln Val 100 105 110

Pro Arg Lys Gly Val Lys Ala Lys Glu Val Phe Gln Arg Phe Leu Glu 115 120 125

Ser Gln Met Val lle Glu Glu Ser lle Leu Gln Ser Asp Lys Ala Leu 130 135 140

Thr Asp Arg Glu Lys Ala Val Ala Val Asp Arg Ala Lys Lys Glu Ala 145 150 155 160

Ala	Glu	Lys	Glu	Gln	Glu	Leu	Leu	Lys	Gln	Lys	Leu	Gln	Glu	Gln	Gln
				165					170					175	
Gln	Gln	Met	Glu	Ala	Gln	Val	Lys	Ser	Arg	Lys	Glu	Asn	Ile	Ala	Gln
			180					185					190		
Leu	Lys	Glu	Lys	Leu	Gln	Met	Glu	Arg	Glu	His	Leu	Leu	Arg	Glu	Gln
		195					200					205			
He	Met	Met	Leu	Glu	His	Thr	Gln	Lys	Val	Gln	Asn	Лsp	Trp	Leu	His
	210					215					220				
Glu	Gly	Phe	Lys	Lys	Lys	Tyr	Glu	Glu	Met	Asn	Ala	Glu	He	Ser	Gln
225					230					235					240
Phe	Lys	Arg	Met	He	Asp	Thr	Thr	Lys	Asn	Asp	Asp	Thr	Pro	Trp	Ile
				245					250					255	
Ala	Arg	Thr	Leu	Asp	Asn	Leu	Ala	Asp	Glu	Leu	Thr	Ala	lle	Leu	Ser
			260					265					270		
Ala	Pro	Ala	Lys	Leu	lle	Gly	His	Gly	Val	Lys	Gly	Val	Ser	Ser	Leu
		275					280					285			
Phe	Lys	Lys	His	Lys	Leu	Pro	Phe								
	290					295									

<211> 153

<212> PRT

<213> Homo sapiens

<400> 507

Met Pro Arg Leu Pro Ala His Arg His Ile Pro Ser Phe Asn Pro Arg 10 Gly Trp Glu Pro Asn Lys Leu Trp Gln Thr Asp Val Thr His lle Pro 30 20 25 Glu Phe Gly Lys Leu Arg Tyr Val His Val Ser Ile Asp Leu Ile Ser 40 45 Ala Asn Ala Leu Pro Gly Glu Ser Thr Gly Tyr Val Thr Lys His Leu 55 60 Leu Leu Thr Phe Ala Phe Met Gly Gln Pro Thr Lys Ile Lys Thr Asn 65 70 75 80 Asn Asp Leu Ala Tyr Ala Ser Ser Gln Phe Gln Gln Phe Cys His Met Trp Asn 11e Gln His Ser Thr Gly 11e Pro Tyr Asn Pro Gln Gly Gln Ala Ile Val Glu Cys Ala His Ser Thr Leu Lys Asn Met Leu Lys Lys Gln Lys Arg Gly Val Trp Val Arg Thr Leu Gln His Tyr Trp His Lys Pro Tyr Leu Pro Leu Ile Phe Lys Ile

<210> 4885

<211> 562

<212> PRT

<213> Homo sapiens

<400> 508

Met Leu Asp Thr 11e Ala Arg Ala Leu Gln Asp Leu Gly Arg Gln Val Leu Pro Thr Leu Pro Ser Leu Ser Gln Glu Glu Val Ser Ile Ile Trp Gly Asn Val Ser Glu Phe Val Arg Arg Gln Leu Thr Leu His Lys Gly Val Gln 11e Pro Ala Phe Gly Thr Phe Thr Phe I1e Arg Gln Lys Leu Glu Val Gly Asn Asn Lys Phe lle Leu Ile Gln Arg Pro Val Phe Ile Met Val Gly Lys Leu Val Gln lle His Gly Leu Lys Gln Asn Lys Arg Pro Gly Thr Val Asp Ser Val Leu Ser Ser Arg Glu Ala Leu Arg Lys Trp Pro Ser Ser Val Leu Ala Phe Pro Arg Ile Glu Leu Lys Glu Met

Glu Asn Lys Leu Pro Met Glu Thr Leu Val Glu Glu Cys Gly Glu Asn

	130					135					140				
Arg	Glu	Arg	Lys	Cys	Lys	Leu	Lys	Asp	Gln	Ser	Asp	Lys	Glu	Glu	Gly
145					150					155					160
Thr	Arg	Asp	lle	Ser	Ser	Pro	Lys	Arg	Leu	Arg	Asp	۸rg	G1n	Ala	Leu
				165					170					175	
Phe	Pro	Λla	Lys	Val	Thr	Asn	Val	Ser	Leu	Leu	Glu	Lys	Phe	Glu	Arg
			180					185					190		
Ser	Glu	Ser	Gly	Gly	Lys	He	Met	Thr	Pro	Glu	Ser	Leu	Ser	Tyr	Pro
		195					200					205			
Ser	Cys	Leu	Lys	His	Asp	Ser	Glu	Met	Lys	Pro	Gln	Thr	Ser	Pro	Ala
	210					215					220				
Cys	Gln	Asp	His	Asn	Lys	Ala	Gly	Gln	Glu	Met	Cys	Tyr	Val	Cys	Leu
225					230					235					240
Gln	Arg.	Ala	Gln	Arg	Asn	Ser	Leu	Leu	Tyr	Tyr	Ser	Glu	Glu	Arg	Λrg
				245					250					255	
Arg	Glu	lle		Asp	Glu	Arg	Leu	lle	Gln	Gln	Tyr	Gln		Leu	Lys
			260					265					270		
Asp	Gln		Ala	Leu	Phe	Arg		Gln	Met	Lys	Ser		Ala	Thr	Arg
		275					280				۵,	285			
Glu		Asn	GIn	Lys	Asn		Ala	Tyr	Asn	Leu		Val	Ala	Glu	Ala
7.1	290		11.			295		10	C1	131	300 T	,	C	DI	
	Arg	Asn	HIS	Lys		61u	Lys	Pro	61u		lyr	Lys	Ser	Phe	
305	Λ	1	Δ	D	310	C	D	A 1 .	1	315	A 1	1	I	C1	320
rne	ASP	Lys	Arg	325	Leu	ser	110	ма	330	ASH	MIA	Leu	Lys	Gln 335	Gru
Glu	Tyr	Sor	Ara		Lou	Lou	Lve	Gln		Asn	Acn	Ara	Gla	Glu	Acn
Giu	1 9 1	361	340	561	Leu	Leu	Lys	345	nic t	пър	nsn	AI g	350	014	non .
Glu	Πe	Lvs		Arø	Gln	Tvr	Arø		Leu	Met	Asn	Arø		Glu	Gln
010	110	355	0.111	8	0111	• , 4	360	010	200	0	пор	365	13.00	010	0111
Val	G1n		Thr	Glu	Glu	Leu		Ala	Gln	Arg	Ala		Phe	Leu	Lvs
	370					375					380	•			
Asp		Met	Glu	Glu	Thr		Cys	Tyr	Lys	Arg		Leu	Asp	Ala	Gln
385	-				390		-	-	-	395			•		400
	Lys	Asn	Lys	Pro	Ser	Arg	Leu	Pro	Pro	Phe	G] u	Pro	Asp	Ser	Ser
				405					410					415	
Glu	Pro	He	Phe	Gly	Lys	Asn	Glu	Gly	Glu	Leu	Met	Val	Glu	Lys	Gln

425 430 Lys Arg Glu Gln Asn Tyr Met Lys His Gln Leu Glu Ala Ala Asn 440 445 His Lys Arg Lys Ala Ile Leu His Gln Leu Val Asp Gln Arg Arg Asp 450 455 460 Leu Gln Met Leu Gln Arg Thr Gln Arg Glu His Leu Ala Asp Arg Thr 470 475 480 Ala Glu Leu Glu Arg Val Asn Arg Val Asn Gln Cys Leu Gln Glu Asp 485 490 Trp Glu Arg Ser Ala Ala Met Lys Lys Gln Arg Asp Leu Glu Asp Lys 500 505 510 Ala Phe Glu Arg Ala Ser Asp Lys Leu Phe Leu Leu Asp Gln Cys Glu 520 Lys Tyr Arg Arg Cys Lys Gln Cys Gln Arg Arg Thr Ser Asn Val Gly 530 535 540 Glu Ser Asn Leu Trp Pro Leu Asn Lys Phe Leu Pro Gly Ser Arg Leu 550 555 560 545 Leu Val

<210> 4886

<211> 385

<212> PRT

<213> Homo sapiens

<400> 509

Met Pro Lys Lys Leu Leu Leu Pro Pro Pro Ser Ala Ser Ser Ala

1 5 10 15

Phe Arg Val Pro Arg Ala Arg Pro Val Pro Pro Pro Ala Met Asn Ala 20 25 30

Ala Arg Thr Gly Tyr Arg Val Phe Ser Ala Asn Ser Thr Ala Ala Cys 35 40 45

Thr Glu Leu Ala Lys Arg lle Thr Glu Arg Leu Gly Ala Glu Leu Gly 50 55 60

Lys Ser Val Val Tyr Gln Glu Thr Asn Gly Glu Thr Arg Val Glu lle

65					70					75					80
Lys	Glu	Ser	Val	Arg	Gly	Gln	Asp	Πe	Phe	Пe	11e	Gln	Thr	lle	Pro
				85					90					95	
Arg	Asp	Val	Asn	Thr	Ala	Val	Met	Glu	Leu	Leu	He	Met	Ala	Tyr	Лlа
			100					105					110		
Leu	Lys	Thr	Ala	Cys	Ala	Arg	Asn	Пе	11e	Gly	Val	Пe	Pro	Tyr	Phe
		115					120					125			
Pro	Tyr	Ser	Lys	Gln	Ser	Lys	Met	Arg	Lys	Arg	Gly	Ser	He	Val	Cys
	130					135					140				
Lys	Leu	Leu	Ala	Ser	Met	Leu	Ala	Lys	Ala	Gly	Leu	Thr	His	Ile	Ile
145					150					155					160
Thr	Met	Asp	Leu	His	Gln	Lys	Glu	He	Gln	Gly	Phe	Phe	Ser	Phe	Pro
				165					170					175	
Val	Лsp	Asn	Leu	Arg	Ala	Ser	Pro	Phe	Leu	Leu	Gln	Tyr	lle	Gln	Glu
			180					185					190		
Glu	Пе	Pro	Asn	Tyr	Arg	Asn	Ala	Val	lle	Va]	Ala	Lys	Ser	Pro	Asp
		195					200					205			
Ala	Ala	Lys	Arg	Ala	Gln	Ser	Tyr	Ala	Glu	Arg	Leu	Arg	Leu	Gly	Leu
	210					215					220				
Ala	Val	He	His	Gly	Glu	Ala	Gln	Cys	Thr	Glu	Leu	Asp	Met	Asp	Asp
225					230					235					240
Gly	Arg	His	Ser	Pro	Pro	Met	Val	Lys	Asn	Ala	Thr	Val	His	Pro	Gly
				245					250					255	
Leu	Glu	Leu		Leu	Met	Met	Ala		Glu	Lys	Pro	Pro		Thr	Val
			260					265					270		
Val	Gly		Val	Gly	Gly	Arg			lle	Пe	Val		Tyr	lle	He
		275					280					285			
Asp		Val	Glu	Ser	Phe		Ala	Ala	Ala	Glu		Leu	Lys	Glu	Arg
	290		,			295			 .		300		,		
	Ala	Tyr	Lys	11e	Tyr	Val	Met	Ala	Thr		Gly	ile	Leu	Ser	
305	. 1	.			310	01	61	0	C	315		C.1	17 1	17 1	320
61u	Ala	Pro	Arg		He	61u	Glu	Ser		Val	Asp	Glu	Val		Val
TL :	Λ	Т1	W. I	325	11.2	C1.	V. 1	C1	330	1 .	C1	C	D.	335	7.7
ınr	Asn	inr		rro	His	Glu	vai		Lys	Leu	ыы	Cys		Lys	116
Lvc	The	Vol	340	11.	Sor	Lou	116	345	Son	61	A1.	116	350	A 22.00	11.

360 365 355 His Asn Gly Glu Ser Met Ala Tyr Leu Phe Arg Asn lle Thr Val Asp 375 380 Asp 385 <210> 4887 <211> 621 <212> PRT <213> Homo sapiens <400> 510 Met Asp Leu Leu Thr Arg Glu Asn Val Ala Leu Lys Val Glu Ser Ala 1 Gln Gln Pro Lys Gln Val Leu Lys Met Glu Val Ala Val Leu Lys Lys 25 Leu Gln Gly Lys Asp His Val Cys Arg Phe Ile Gly Cys Gly Arg Asn 40 45 Glu Lys Phe Asn Tyr Val Val Met Gln Leu Gln Gly Arg Asn Leu Ala 50 60 55 Asp Leu Arg Arg Ser Gln Pro Arg Gly Thr Phe Thr Leu Ser Thr Thr 75 Leu Arg Leu Gly Lys Gln 11e Leu Glu Ser 11e Glu Ala 11e His Ser 90 85 Val Gly Phe Leu His Arg Asp Ile Lys Pro Ser Asn Phe Ala Met Gly 100 105 Arg Leu Pro Ser Thr Tyr Arg Lys Cys Tyr Met Leu Asp Phe Gly Leu 120 125 Ala Arg Gln Tyr Thr Asn Thr Thr Gly Asp Val Arg Pro Pro Arg Asn 130 135 140 Val Ala Gly Pho Arg Gly Thr Val Arg Tyr Ala Ser Val Asn Ala His 150 155 Lys Asn Arg Glu Met Gly Arg His Asp Asp Leu Trp Ser Leu Phe Tyr 165 170

Met Leu Val Glu Phe Ala Val Gly Gln Leu Pro Trp Arg Lys Ile Lys

			180					185					190		
Asp	Lys	Glu	Gln	Val	Gly	Met	He	Lys	Glu	Lys	Tyr	Glu	His	Arg	Met
		195					200					205			
Leu	Leu	Lys	His	Met	Pro	Ser	G1u	Phe	His	Leu	Phe	Leu	Asp	His	He
	210					215					220				
Λla	Ser	Leu	Asp	Tyr	Phe	Thr	Lys	Pro	Asp	Tyr	G1n	Leu	He	Met	Ser
225					230					235					240
Val	Phe	Glu	Asn	Ser	Met	Lys	Glu	Arg	Gly	11e	Ala	Glu	Asn	Glu	Ala
				245					250					255	
Phe	Asp	Trp	Glu	Lys	Λla	Gly	Thr	Лѕр	Ala	Leu	Leu	Ser	Thr	Ser	Thr
			260					265					270		
Ser	Thr	Pro	Pro	Gln	Gln	Asn	Thr	Arg	Gln	Thr	Ala	Ala	Met	Phe	Gly
		275					280					285			
Val	Val	Asn	Val	Thr	Pro	Val	Pro	Gly	Asp	Leu	Leu	Arg	Glu	Asn	Thr
	290					295					300				
G1u	Asp	Val	Leu	Gln	Gly	Glu	His	Leu	Ser	Asp	Gln	Glu	Asn	Ala	Pro
305					310					315					320
Pro	Ile	Leu	Pro	Gly	Arg	Pro	Ser	Glu	Gly	Leu	Gly	Pro	Ser	Pro	His
				325					330			•		335	
Leu	Val	Pro	His	Pro	Gly	Gly	Pro	Glu	Ala	Glu	Val	Trp	Glu	Gly	Thr
			340					345					350		
Asp	Val	Asn	Arg	Asn	Lys	Leu	Arg	Пе	Asn	He	Gly	Lys	Ser	Pro	Cys
		355					360					365			
Val	Glu	Glu	Glu	Gln	Ser	Arg	G] y	Met	Gly	Val	Pro	Ser	Ser	Pro	Val
	370					375					380				
Arg	Ala	Pro	Pro	Asp	Ser	Pro	Thr	Thr	Pro	Va]	Arg	Ser	Leu	Arg	Tyr
385					390					395					400
Arg	Arg	Val	Asn	Ser	Pro	Glu	Ser	G}u	Arg	Leu	Ser	Thr	Ala	Asp	Gly
				405					410					415	
Arg	Val	Glu		Pro	Głu	Arg	Arg		Arg	Met	Asp	Leu		Gly	Ser
			420					425					430		
Pro	Ser		Gln	Ala	Cys	Ser	Ser	Gln	Pro	Ala	Gln		Leu	Ser	Val
		435					440					445	. —		
Asp		Gly	His	Ala	Asp		Gln	Ala	Ser	Gly		Met	Asp	Val	Ser
	450					455					460				

Ala Ser Val Glu Glu Ala Leu Ser Asn Ala Phe Arg Ser Val Pro Leu Ala Glu Glu Glu Asp Phe Asp Ser Lvs Glu Trp Val lle lle Asp Lys Glu Thr Glu Leu Lys Asp Phe Pro Pro Gly Ala Glu Pro Ser Thr Ser Gly Thr Thr Asp Glu Glu Pro Glu Glu Leu Arg Pro Leu Pro Glu Glu Gly Glu Glu Arg Arg Leu Gly Ala Glu Pro Phe Pro Leu Thr Pro Ala Leu Gly Thr Pro Pro Ser Thr Glu Arg Val Gly Pro His Arg Pro Thr Glu Thr Val Gly Gly Gly Gln Thr Leu Gly Ala Leu Pro Pro Ala Val Gln Pro Pro Ala Thr Thr Gly Val Leu Arg Val Leu Leu Leu His Ala Gly Asp Gly Ala Leu Pro Ser Pro Arg Arg Arg Leu Leu Gly Leu Leu Arg Phe Pro His Ser Ala Gln Pro Leu Gly

<210> 4888

<211> 267

<212> PRT

<213> Homo sapiens

<400> 511

Met Phe Arg Arg Lys Ala Glu Gly Leu Asp Leu Ala Ser Cys Val Arg l Ser Leu Asp Val Leu Val Leu Asp Glu Ala Asp Arg Leu Leu Asp Met Gly Phe Glu Ala Ser Ile Asn Thr Ile Leu Glu Phe Leu Pro Lys Gln Arg Arg Thr Gly Leu Phe Ser Ala Thr Gln Thr Gln Glu Val Glu Asn

Leu Val Arg Ala Gly Leu Arg Asn Pro Val Arg Val Ser Val Lys Glu Lys Gly Val Ala Ala Ser Ser Ala Gln Lys Thr Pro Ser Arg Leu Glu Asn Tyr Tyr Met Val Cys Lys Ala Asp Glu Lys Phe Asn Gln Leu Val His Phe Leu Arg Asn His Lys Gln Glu Lys His Leu Val Phe Phe Ser Thr Cys Ala Cys Val Glu Tyr Tyr Gly Lys Ala Leu Glu Val Leu Val Lys Gly Val Lys Ile Met Cys Ile His Gly Lys Met Lys Tyr Lys Arg Asn Lys lle Phe Met Glu Phe Arg Lys Leu Gln Ser Gly lle Leu Val Cys Thr Asp Val Met Ala Arg Gly lle Asp lle Pro Glu Val Asn Trp Val Leu Gln Tyr Asp Pro Pro Ser Asn Ala Ser Ala Phe Val His Arg Cys Gly Arg Thr Ala Arg Ile Gly His Gly Gly Ser Ala Leu Val Phe Leu Leu Pro Met Glu Glu Ser Tyr lle Asn Phe Leu Ala lle Ser Gln Lys Val Ser Cys Arg Pro Phe Ser Asp Arg Met Pro Ser Asp Gly Val Ala Gly Lys Val Leu Gln His Val Val Ser Asn

<210> 4889

<211> 700

<212> PRT

<213> Homo sapiens

<400> 512

Met Leu Ser Asp Asp His Val Asn Glu Ile Ile Ile Gln Lys Leu Ile

1 5 10 15

Ala	Ser	Leu	He	Pro	Met	Thr	Ser	Arg	Asp	Arg	He	Lys	Ala	lle	Arg
			20					25					30		
Asn	Gln	Pro	Arg	Thr	Met	Glu	Glu	Lys	۸rg	Asn	Leu	Ser	Arg	Gly	Gly
		35					40					45			
Leu	Thr	lle	Thr	Thr	Glu	Asp	Glu	Gly	Arg	Ala	Lys	Ala	His	Leu	Thr
	50					55					60				
Trp	Trp	Lys	He	Val	Asp	Lys	Glu	Lys	Ser	Lys	Gln	Thr	His	Arg	lle
65					70					75					80
Leu	Gln	Leu	Asn	Cys	Cys	Ile	Gln	Cys	Leu	Asn	Ser	lle	Ser	Arg	Ala
				85					90					95	
Tyr	Arg	Arg	Ser	Lys	Asn	Ser	Leu	Ser	Glu	Ile	Leu	Asn	Ser	11e	Ser
			100					105					110		
Leu	Trp	Gln	Lys	Thr	Leu	Lys		He	Gly	G1 y	Lys		Gly	Thr	Ser
		115					120					125			
Va]		Ser	Tyr	Phe	Asn	Phe	Leu	Arg	Trp	Leu	Leu	Lys	Phe	Asn]]e
	130					135					140				
	Ser	Phe	lle	Leu		Phe	Ser	Phe	lle		Ile	Pro	Gln	Phe	
145					150					155					160
Val	Ala _.	Lys	Lys		Thr	Leu	GIn	Phe		Gly	Leu	Glu	Phe		Thr
0.1		0.1	m.	165			mı	., .	170	m	ar.	0.1	DI	175	T)
Gly	Val	Gly		Phe	Arg	Asp	Ihr		Met	lyr	lyr	Gly		lyr	lhr
	C	TI	180	61	11.	61		185	C1	4.7	C	T.	190		C I
Asn	Ser		11e	GIn	HIS	G1 y		Ser	61 y	Ala	261.		Asn	мет	GIn
1	410	195	Tla	Dha	Tha	11.	200	A16	Cva	Lau	The	205	Cua	Dlag	Dho
Leu	210	1 9 1	116	rne	1111	11e 215	Oly	Ма	Cys	Leu	220	1111	Cys	rne	THE
Sor		Lou	Pho	Sor	Mat	Ala	Lve	Tyr	Pho	Ara		Acn	Pho	مال	Aen
225	Leu	Leu	THE	361	230	ма	rys	1 y 1	1116	235	лэн	изп	1116	116	240
	His	Tle	Tvr	Ser		Gly	He	Thr	lvs		He	Phe	Cvs	Trn	
,,,	1113	110	1,1	245	019	0.1 9	110	****	250	150.0	110	1110	0,0	255	11
Phe	Thr	Val	Thr		Glu	Lys	Ala	Val		Leu	Lvs	Gln	Lvs		Leu
			260			-,-		265	_,.				270		
Ser	Thr	Glu		Arg	Glu	Asn	Leu		Glu	Leu	Arg	Gln		Asn	Ser
		275		9			280				0	285			
Lys	Leu		Phe	Asn	Gln	Leu		Thr	Arg	Phe	Ser		Tyr	Met	Val
-	290					295			J		300		•		

Ala	Trp	Val	Val	Ser	Thr	Gly	Val	Ala	He	Ala	Cys	Cys	Ala	Ala	Val
305					310					315					320
Tyr	Tyr	Leu	Ala	Glu	Tyr	Asn	Leu	Glu	Phe	Leu	Lys	Thr	His	Ser	Asn
				325					330					335	
Pro	Gly	Ala	Val	Leu	Leu	Leu	Pro	Phe	Val	Val	Ser	Cys	11e	Asn	Leu
			340					345					350		
Ala	Val	Pro	Cys	He	Tyr	Ser	Met	Phe	Arg	Leu	Val	Glu	Arg	Tyr	Glu
		355					360					365			
Met	Pro	Arg	His	Glu	Val	Tyr	Val	Leu	Leu	Пe	Arg	Asn	Ile	Phe	Leu
	370					375					380				
Lys	Ile	Ser	He	Ile	Gly	lle	Leu	Cys	Tyr	Tyr	Trp	Leu	Asn	Thr	Val
385					390					395					400
Ala	Leu	Ser	Gly	Glu	Glu	Cys	Trp	Gly	Thr	Leu	He	Gly	Gln	Asp	11e
				405					410					415	
Tyr	Arg	Leu	Leu	Leu	Met	Лsp	Phe	Val	Phe	Ser	Leu	Val	Asn	Ser	Phe
			420					425					430		
Leu	Gly	Glu	Phe	Leu	Arg	Arg	Ile	lle	Gly	Met	Gln	Leu	Ile	Thr	Ser
		435					440					445			
Leu	Gly	Leu	Gln	Glu	Phe	Asp	lle	Ala	Arg	Asn	Val	Leu	Glu	Leu	He
	450					455					460				
Tyr	Ala	Gln	Thr	Leu	Val	Trp	lle	Gly	He	Phe	Phe	Arg	Pro	Leu	Leu
465					470					475					480
Pro	Phe	Ile	Gln	Met	lle	Met	Leu	Phe	He	Met	Phe	Tyr	Ser	Lys	Asn
				485					490					495	
Πle	Ser	Leu	Met	Met	Asn	Phe	Gln	Pro	Pro	Ser	Lys	Ala	Trp	Arg	Ala
			500					505					510		
Ser	Gln	Met	Met	Thr	Phe	Phe	lle	Phe	Leu	Leu	Phe	Phe	Pro	Ser	Phe
		515					520					525			
Thr	Gly	Val	Leu	Cys	Thr	Leu	Ala	He	Thr	He	Trp	Arg	Leu	Lys	Pro
	530					535					540				
Ser	Ala	Asp	Cys	Gly	Pro	Phe	Arg	Gly	Leu	Pro	Leu	Phe	lle	His	Ser
545					550					555					560
He	Tyr	Ser	Trp	lle	Asp	Thr	Leu	Ser	Thr	Arg	Pro	Gly	Tyr	Leu	Trp
				565					570					575	
Val	Val	Trp	He	Tyr	Arg	Asn	Leu	He	Gly	Ser	Val	His	Phe	Phe	Phe
			580					585					590		

lle Leu Thr Leu Ile Val Leu Ile Ile Thr Tyr Leu Tyr Trp Gln Ile 600 605 Thr Glu Gly Arg Lys Ile Met Ile Arg Leu Leu His Glu Gln Ile Ile 610 615 620 Asn Glu Gly Lys Asp Lys Met Phe Leu Ile Glu Lys Leu Ile Lys Leu 630 635 640 625 Gln Asp Met Glu Lys Lys Ala Asn Pro Ser Ser Leu Val Leu Glu Arg 645 650 Arg Glu Val Glu Gln Gln Gly Phe Cys Ile Trp Gly Asn Met Met Ala 660 665 670 Val Leu Thr Cys Asp Leu Glu Asp Gln Phe Lys Lys Ala Ile Gln Gly 680 685 Pro Asp Asp Ser Phe Gly Asn Gln Thr Pro 11e Lys 690 695

<210> 4890

<211> 174

<212> PRT

<213> Homo sapiens

<400> 513

Met Ile Ser Asn Leu Ser Trp Glu Leu Pro Gly Ser Leu Pro Leu Ile 1 5 10 15

Ser Val Pro Tyr Ser Met His Cys Cys Thr Leu Gly Phe Leu Ser Cys 20 25 30

Ser Leu Phe Leu His Met Ser Phe Glu Leu Lys Leu Leu Leu Leu Leu 35 40 45

Leu Trp Leu Ala Ala Ser Cys Ser Leu Phe Leu His Ser His Ala Trp 50 55 60

Leu Ser Glu Cys Leu Ile Val Arg Leu Tyr Leu Gly Pro Leu Asp Ser 65 70 75 80

Arg Pro Gly Val Leu Lys Glu Pro Lys Leu Met Gly Ala 11e Ser Phe 85 90 95

Phe lle Phe Phe Phe Thr Leu Leu Val Leu Ala Arg Gln Asn Glu Tyr 100 105 110 Tyr Cys Arg Leu Asp Phe Leu Trp Lys Lys Leu Arg Gln Glu Arg Glu Glu Thr Glu Thr Met Glu Asn Leu Thr Arg Leu Leu Glu Asn Val Leu Pro Ala His Val Ala Pro Gln Phe Ile Gly Gln Asn Arg Arg Asn Glu Ser Pro Val Pro Pro Gly Ser Leu Pro Pro Val Leu

<210> 4891

<211> 546

<212> PRT

<213> Homo sapiens

<400> 514 Met lle His Thr Thr Glu Lys Pro Tyr Arg Cys Asn Glu Ser Gly Lys Ala Phe His Arg Gly Ser Leu Leu Thr Val His Gln Ile Val His Thr Arg Gly Lys Pro Tyr Gln Cys Asp Val Cys Gly Arg Ile Phe Arg Gln Asn Ser Asp Leu Val Asn His Arg Arg Ser His Thr Gly Asp Lys Pro Tyr Ile Cys Asn Glu Cys Gly Lys Ser Phe Ser Lys Ser Ser His Leu Ala Val His Gln Arg Ile His Thr Gly Glu Lys Pro Tyr Lys Cys Asn Arg Cys Gly Lys Cys Phe Ser Gln Ser Ser Ser Leu Ala Thr His Gln Thr Val His Thr Gly Asp Lys Pro Tyr Lys Cys Asn Glu Cys Gly Lys Thr Phe Lys Arg Asn Ser Ser Leu Thr Ala His His Ile Ile His Ala

Gly Lys Lys Pro Tyr Thr Cys Asp Val Cys Gly Lys Val Phe Tyr Gln

Asn	Ser	Gln	Leu	Val	Arg	His	Gln	Пe	lle	His	Thr	Gly	Glu	Thr	Pro
				165					170					175	
Tyr	Lys	Cys	Asn	Glu	Cys	Gly	Lys	Val	Phe	Phe	Gln	Arg	Ser	Arg	Leu
			180					185					190		
Ala	Gly	His	Arg	Arg	He	His	Thr	Gly	Glu	Lys	Pro	Tyr	Lys	Cys	Asn
		195					200					205			
Glu	Cys	Gly	Lys	Val	Phe	Ser	Gln	His	Ser	His	Leu	Ala	Val	His	Gln
	210					215					220				
Arg	Val	His	Thr	Gly	Glu	Lys	Pro	Tyr	Lys	Cys	Asn	Glu	Cys	Gly	Lys
225					230					235					240
Ala	Phe	Asn	Trp	Gly	Ser	Leu	Leu	Thr	Val	His	Gln	Arg	Ile	His	Thr
				245					250					255	
Gly	Glu	Lys	Pro	Tyr	Lys	Cys	Asn	Val	Cys	Gly	Lys	Val	Phe	Asn	Tyr
			260					265					270		
Gly	Gly	Tyr	Leu	Ser	Val	His	Met	Arg	Cys	His	Thr	Gly	Glu	Lys	Pro
		275					280					285			
Leu	His	Cys	Asn	Lys	Cys	Gly	Met	Val	Phe	Thr	Tyr	Tyr	Ser	Cys	Leu
	290					295					300				
Ala	Arg	His	Gln	Arg	Met	His	Thr	Gly	Glu	Lys	Pro	Tyr	Lys	Cys	Asn
305					310					315					320
Val	Cys	Gly	Lys	Val	Phe	Ile	Asp	Ser	Gly	Asn	Leu	Ser	He	His	Arg
				325					330					335	
Arg	Ser	His	Thr	Gly	Glu	Lys	Pro	Phe	Gln	Cys	Asn	Glu	Cys	Gly	Lys
			340					345					350		
Val	Phe	Ser	Tyr	Tyr	Ser	Cys	Leu	Ala	Arg	His	Arg	Lys	lle	His	Thr
		355					360					365			
Gly	Glu	Lys	Pro	Tyr	Lys	Cys	Asn	Asp	Cys	G] y	Lys	Ala	Tyr	Thr	Gln
	370					375					380				
Arg	Ser	Ser	Leu	Thr	Lys	His	Leu	Val	lle	His	Thr	Gly	Glu	Asn	Pro
385					390					395					400
Tyr	llis	Cys	Asn	Glu	Phe	Gly	Glu	Ala	Phe	lle	Gln	Ser	Ser	Lys	Leu
				405					410					415	
Ala	Arg	Tyr	His	Arg	Asn	Pro	Thr	Gly	Glu	Lys	Pro	His	Lys	Cys	Ser
			420					425					430		
Glu	Cys	Gly	Arg	Thr	Phe	Ser	His	Lys	Thr	Ser	Leu	Val	Tyr	His	Gln
		435					440					445			

Arg Arg His Thr Gly Glu Met Pro Tyr Lys Cys Ile Glu Cys Gly Lys 455 Val Phe Asn Ser Thr Thr Leu Ala Arg His Arg Arg Ile His Thr 465 470 475 480 Gly Glu Lys Pro Tyr Lys Cys Asn Glu Cys Gly Lys Val Phe Arg Tyr 485 490 Arg Ser Gly Leu Ala Arg His Trp Ser Ile His Thr Gly Glu Lys Pro 505 Tyr Lys Cys Asn Glu Cys Gly Lys Ala Phe Arg Val Arg Ser Ile Leu 515 520 525 Leu Asn His Gln Met Met His Thr Gly Glu Lys Pro Tyr Lys Cys Asn 530 535 540 Glu Cys 545

<210> 4892

<211> 113

<212> PRT

<213> Homo sapiens

<400> 515

Met Met His Leu Gln Pro Ala Trp Ser Ser Phe Gly Gln Ser Gly Pro

1 5 10 15

Ile Leu Met Arg Ser Gly Gly Met Gln Leu Leu Leu Leu Pro His Gly
20 25 30

Ser Val Arg Pro Asn Gln Pro Ala Leu Ser Pro Leu Pro Ala Val Tyr 35 40 45

Gln Val Ala Leu Glu Arg Gly His Pro Glu Gly Thr Thr Cys Ala Ala 50 55 60

Val Gly Gly Lys lle Asn Arg Pro Arg Thr Pro Gly Gln Ala Gly
65 70 75 80

Ala Thr Glu Glu Val Ser Ser Lys Gly Gly Lys Leu Ser Gly Leu Phe
85 90 95

Leu Thr Ile Leu Pro Val Pro Arg Pro Trp Thr Ala Gly Gln Pro Gln
100 105 110

Ser

<210> 4893

<211> 129

<212> PRT

<213> Homo sapiens

<400> 516

Met Met Thr Asp Leu Lys Gln Ser His Ser Val Arg Leu Asn Asp Gly

1 5 10 15

Pro Phe Met Pro Val Leu Gly Phe Gly Thr Tyr Ala Pro Asp His Thr
20 25 30

Pro Lys Ser Gln Ala Ala Glu Ala Thr Lys Val Ala Ile Asp Val Gly
35 40 45

Phe Arg His Ile Asp Ser Ala Tyr Leu Tyr Gln Asn Glu Glu Glu Val
50 55 60

Gly Gln Ala Ile Trp Glu Lys Ile Ala Asp Gly Thr Val Lys Arg Glu
65 70 75 80

Glu Ile Phe Tyr Thr Ile Lys Leu Trp Ala Thr Phe Phe Arg Ala Glu 85 90 95

Leu Val His Pro Ala Leu Glu Arg Ser Leu Lys Lys Leu Gly Pro Asp 100 105 110

Tyr Val Asp Leu Phe Ile Ile His Val Pro Phe Ala Met Lys Gly Ser 115 120 125

Ser

<210> 4894

<211> 105

<212> PRT

<213> Homo sapiens

<400> 517

Met Cys Pro Val Cys Gly Arg Ala Leu Ser Ser Pro Gly Ser Leu Gly 10 Arg His Leu Leu IIe His Ser Glu Asp Gln Arg Ser Asn Cys Ala Val 25 Cys Gly Ala Arg Phe Thr Ser His Ala Thr Phe Asn Arg Ser Ala Gly 35 40 45 His Leu Ser Leu Leu Cys Trp Glu Gln lle Thr Gly Cys Glu Phe Cys 55 Phe Leu Lys Ser Lys Arg Arg Leu Glu Ile Gly Asn Ala Lys Ala Ala 65 70 75 Asp Ala Cys Ser Leu Glu Val Gln Val Arg Arg Leu Gly Asn Met Pro 85 90 95 lle Pro Ser His Ser Pro Lys Gly Lys 100

<210> 4895

<211> 646

<212> PRT

<213> Homo sapiens

<400> 518

Met Phe Leu Ser Ile Ser Thr Arg Leu Pro Ser Gln Tyr Ile Tyr Gly
1 5 10 15

Phe Gly Glu Thr Glu His Thr Thr Phe Arg Arg Asn Met Asn Trp Asn 20 25 30

Thr Trp Gly Met Phe Ala His Asp Glu Pro Pro Ala Tyr Lys Lys Asn 35 40 45

Ser Tyr Gly Val His Pro Tyr Tyr Met Ala Leu Glu Glu Asp Gly Ser 50 55 60

Ala His Gly Val Leu Leu Leu Asn Ser Asn Ala Met Asp Val Thr Leu 65 70 75 80

Gln Pro Thr Pro Ala Leu Thr Tyr Arg Thr Thr Gly Gly lle Leu Asp 85 90 95

Phe Tyr Ile Val Leu Gly Pro Thr Pro Glu Leu Val Thr Gln Gln Tyr 100 105 110

Thr	Glu	Leu	He	Gly	Arg	Pro	Ala	Met	lle	Pro	Tyr	Trp	Ala	Leu	Gly
		115					120					125			
Phe	His	Leu	Ser	Arg	Tyr	Gly	Tyr	Gln	Asn	Asp	Ala	Glu	Пe	Ser	Ser
	130					135					140				
Leu	Tyr	Asp	Ala	Met	Val	Ala	Ala	Gln	He	Pro	Tyr	Asp	Val	Gln	His
145					150					155					160
Val	Asp	lle	Asp	Tyr	Met	Asn	Arg	Lys	Leu	Asp	Phe	Thr	Leu	Ser	Ala
				165					170					175	
Asn	Phe	Gln	Asn	Leu	Ser	Leu	Leu	lle	Glu	Gln	Met	Lys	Lys	Asn	Gly
			180					185					190		
Met	Arg	Phe	He	Leu	Ile	Leu	Asp	Pro	Ala	Ile	Ser	Gly	Asn	Glu	Thr
		195					200					205			
Gln	Tyr	Leu	Pro	Phe	lle	Arg	Gly	Gln	Glu	Asn	Asn	Val	Phe	He	Lys
	210					215					220				
Trp	Pro	Asp	Thr	Asn	Asp	lle	Val	Trp	Gly	Lys	Val	Trp	Pro	Asp	Leu
225					230					235					240
Pro	Asn	Val	lle	Val	Asp	Gly	Ser	Leu	Asp	His	Glu	Thr	Gln	Val	Lys
				245					250					255	
Leu	Tyr	Arg	Ala	Tyr	Va]	Ala	Phe	Pro	Asp	Phe	Phe	Arg	Asn	Ser	Thr
			260					265					270		
Ala	Ala	Trp	Trp	Lys	Lys	Glu	lle	Glu	Glu	Leu	Tyr	Ala	Asn	Pro	Arg
		275					280					285			
Glu	Pro	Glu	Lys	Ser	Leu	Lys	Phe	Asp	Gly	Leu	Trp	He	Asp	Met	Asn
	290					295					300				
Glu	Pro	Ser	Asn	Phe	Val	Asp	Gly	Ser	Val	Arg	Gly	Cys	Ser	Asn	Glu
305					310					315					320
Met	Leu	Asn	Asn	Pro.	Pro	Tyr	Met	Pro	Tyr	Leu	Glu	Ser	Arg	Asp	Lys
				325					330					335	
Gly	Leu	Ser	Ser	Lys	Thr	Leu	Cys	Met	Glu	Ser	Gln	Gln	lle	Leu	Pro
			340					345					350		
Asp	Ser	Ser	Pro	Val	Glu	His	Tyr	Asn	Val	His	Asn	Leu	Tyr	Gly	Trp
		355					360					365			
Ser		Thr	Arg	Pro	Thr	-	Glu	Ala	Val	Gln		Val	Thr	Gly	Gln
	370					375					380				
	Gly	Val	11e	He		Arg	Ser	Thr	Phe		Ser	Ser	G] y	Arg	
385					390					395					400

Gly	Gly	His	Arg		Gly	Asn	Asn	Thr	Ala	Ala	Trp	Asp	Gln	Leu	Gly
				405					410					415	
Lys	Ser	He	He	G1 y	Met	Met	Glu	Phe	Ser	Leu	Phe	Gly	He	Pro	Tyr
			420					425					430		
Thr	Gly	Ala	Asp	lle	Cys	Gly	Phe	Phe	Gly	Asp	Ala	Glu	Tyr	Glu	Met
		435					440					445			
Cys	Val	Arg	Trp	Met	Gln	Leu	Gly	Ala	Phe	Tyr	Pro	Phe	Ser	Arg	Asn
	450					455					460				
His	Asn	Asn	He	Gly	Thr	Arg	Arg	Gln	Asp	Pro	Val	Ala	Trp	Asn	Ser
465					470					475					480
Thr	Phe	Glu	Met	Leu	Ser	Arg	Lys	Val	Leu	Glu	Thr	Arg	Tyr	Thr	Leu
				485					490					495	
Leu	Pro	Tyr	Leu	Tyr	Thr	Leu	Met	His	Lys	Ala	His	Va]	Glu	Gly	Ser
			500					505					510		
Thr	Val	Val	Arg	Pro	Leu	Leu	His	Glu	Phe	Thr	Asp	Asp	Arg	Thr	Thr
		515					520					525			
Trp	Asp	lle	Asp	Arg	Gln	Phe	Met	Leu	Gly	Pro	Ala	Ile	Leu	Ile	Ser
	530					535					540				
Pro	Val	Leu	Glu	Thr	Ser	Thr	Phe	Glu	He	Ser	Ala	Tyr	Phe	Pro	Arg
545					550					555					560
Ala	Arg	Trp	Tyr	Asp	Tyr	Ser	Thr	Gly	Thr	Ser	Ser	Thr	Ser	Thr	G1 y
				565					570					575	
Gln	Arg	Lys	lle	Leu	Lys	Ala	Pro	Leu	Asp	His	lle	Asn	Leu	His	Val
			580					585					590		
Arg	Gly	Gly	Tyr	lle	Leu	Pro	Trp	Gln	Glu	Pro	Ala	Met	Asn	Thr	His
		595					600					605			
Ser	Ser	Arg	Gln	Asn	Phe	Met	Gly	Leu	lle	Val	Ala	Leu	Asp	Asp	Asn
	610					615					620				
Gly	Thr	Ala	Glu	Gly	Gln	Val	Phe	Trp	Asp	Asp	Gly	Gln	Ser	He	Val
625	•				630					635					640
Phe	Asn	Thr	Thr	Ala	Met										
				645											

<210> 4896

<211> 472

<212	2> PI	T5													
<213	3> He	omo s	sapie	ens											
<400)> 51	19													•
Met	Ala	Ser	Pro	Pro	Arg	Cys	Ser	Pro	Thr	Ala	His	Asp	Arg	Glu	Cys
1				5					10					15	
Lys	Leu	Pro	Pro	Pro	Ser	Ala	Pro	Ala	Ser	Glu	Tyr	Cys	Pro	Gly	Lys
			20					25					30		
Leu	Ser	Trp	Gly	Thr	Met	Ala	Arg	Ala	Leu	Gly	Arg	Phe	Lys	Leu	Ser
		35					40					45			
He	Pro	His	Thr	His	Leu	Leu	Ala	Thr	Leu	Asp	Pro	Leu	Ala	Leu	Asp
	50					55					60				
Arg	Glu	Pro	Pro	Pro	His	Leu	Leu	Pro	Glu	Lys	His	Gln	Val	Pro	Glu
65					70					75					80
Lys	Leu	He	Trp	Gly	Asp	Gln	Asp	Pro	Leu	Ser	Lys	lle	Pro	Phe	Lys
				85					90					95	
He	Leu	Ser	Gly	His	Glu	His	Ala	Val	Ser	Thr	Cys	His	Phe	Cys	Val
			100					105					110		
Asp	Asp	Thr	Lys	Leu	Leu	Ser	Gly	Ser	Tyr	Asp	Cys	Thr	Val	Lys	Leu
		115					120					125			
Trp	Asp	Pro	Va]	Asp	Gly	Ser	Val	Val	Arg	Asp	Phe	Glu	His	Arg	Pro
	130					135					140				
Lys	Ala	Pro	Val	Val	Glu	Cys	Ser	He	Thr	G] y	Asp	Ser	Ser	Arg	Val
145					150					155					160
He	Ala	Ala	Ser	Tyr	Asp	Lys	Thr	Val	Arg	Ala	Trp	Asp	Leu	Glu	Thr
				165					170					175	
Gly	Lys	Leu	Leu	Trp	Lys	Val	Arg	Tyr	Asp	Thr	Phe	He	Val	Ser	Cys
			180					185					190		
Lys	Phe		Pro	Asp	G] y	Lys		Va1	Val	Ser	Gly		Asp	Va]	Asp
		195					200					205			
llis]]e	Cys	He	Met		Ala	Glu	Asn	lle		Thr	Val	Ser	Val
	210					215					220				

He Lys Asp His His Thr Arg Ser He Thr Ser Cys Cys Phe Asp Pro

Asp Ser Gln Arg Val Ala Ser Val Ser Leu Asp Arg Cys 11e Lys 11e

Trp	Asp	Val	Thr	Ser	Gln	Ala	Thr	Leu	Leu	Thr	He	Thr	Lys	Ala	His
			260					265					270		
Ser	Asn	Ala	lle	Ser	Asn	Cys	Cys	Phe	Thr	Phe	Ser	Gly	His	Phe	Leu
		275					280					285			
Cys	Thr	Ser	Ser	Trp	Asp	Lys	Asn	Leu	Lys	lle	Trp	Asn	Val	His	Thr
	290					295					300				
G1 y	Glu	Phe	Arg	Asn	Arg	Gly	Ala	Cys	Val	Thr	Leu	Met	Gln	Gly	His
305					310					315					320
Glu	Gly	Ser	Val	Ser	Ser	Cys	His	Phe	Ala	Arg	Asp	Ser	Ser	Phe	Leu
				325					330					335	
He	Ser	Gly	Gly	Phe	Asp	Arg	Thr	Val	Λla	He	Trp	Asp	Val	Ala	Glu
			340					345					350		
Gly	Tyr	Arg	Lys	Leu	Ser	Leu	Lys	Gly	llis	Asn	Asp	Trp	Val	Met	Asp
		355					360					365			
Val	Ala	He	Ser	Asn	Asn	Lys	Lys	Trp	Пе	Leu	Ser	Ala	Ser	Lys	Asp
	370					375					380				
Arg	Thr	Met	Arg	Leu	Trp	Asn	lle	Glu	Glu	He	Asp	Glu	Πle	Pro	Leu
385					390					395					400
Val	He	Lys	Tyr	Lys	Lys	Ala	Val	Gly	Leu	Lys	Leu	Lys	Gln	Cys	Glu
				405					410					415	
Arg	Cys	Asp	Arg	Pro	Phe	Ser	lle	Phe	Lys	Ser	Asp	Thr	Ser	Ser	Glu
			420					425					430		
Met	Phe	Thr	Gln	Cys	Va]	Phe	Cys	Arg	lle	Asp	Thr	Arg	Gly	Leu	Pro
		435					440					445			
Ala	Asp	Thr	Ser	Glu	Arg	Glu	Asn	Ser	Pro						
	450					455					460				
Pro	Pro	Arg	Gly	Ser	Lys	Asp	Asp								
465					470										

<210> 4897

<211> 466

<212> PRT

<213> Homo sapiens

<400	> 52	20													
			Glv	Val	Ala	Arg	His	Ser	Glv	Ser	Gln	Asp	Glu	Val	Ser
1				5		0			10					15	
	Glv	Val	Glu		Leu	Asp	Λla	Ala		Ala	Gln	Pro	Ala		Asp
o o	•		20					25					30		•
Arg	۸rg	Ala	Lvs	Glv	Thr	Pro	Lvs	Ser	Ser	Lvs	Pro	Glv	Lys	Lvs	His
	Ŭ	35	•	•			40			•		45	•	•	
Arg	Tyr	Leu	Arg	Leu	Leu	Pro	Glu	Ala	Leu	lle	Arg	Phe	Gly	G1 y	Phe
	50					55					60				
Arg	Lys	Arg	Lys	Lys	Ala	Lys	Ser	Ser	Val	Ser	Lys	Lys	Pro	Gly	Glu
65					70					75					80
/al	Asp	Asp	Ser	Leu	Glu	Gln	Pro	Cys	Gly	Leu	Gly	Cys	Leu	Val	Ser
				85					90					95	
Γhr	Cys	Cys	Glu	Cys	Cys	Asn	Asn	He	Arg	Cys	Phe	Met	He	Phe	Tyr
			100					105					110		
Cys	He	Leu	Leu	Ile	Cys	Gln	Gly	Val	Val	Phe	Gly	Leu	He	Asp	Val
		115					120					125			
Ser	Ile	Gly	Asp	Phe	Gln	Lys	Glu	Tyr	Gln	Leu	Lys	Thr	Ile	Glu	Lys
	130					135					140				
.eu	Ala	Leu	Glu	Lys	Ser	Tyr	Asp	He	Ser	Ser	Gly	Leu	Val	Ala	Пe
145					150					155					160
Phe	lle	Ala	Phe	Tyr	Gly	Asp	Arg	Lys	Lys	Val	He	Trp	Phe	Val	Ala
				165					170					175	
Ser	Ser	Phe	Leu	He	Gly	Leu	G1 y	Ser	Leu	Leu	Cys	Ala	Phe	Pro	Ser
			180					185					190		
He	Asn		Glu	Asn	Lys	Gln			Val	Gly	He		Gly	He	Ala
		195	_				200		_		_	205	_		
		Thr	Ser	Met	He		Tyr	Ala	Leu	Gly		Val	Leu	Gly	Ala
	210	., .	,			215		T)	T.I		220	m		771	0.1
	Leu	Val	Lys	Va.I		GJu	Asn	ihr	lhr		Ala	Thr	Lys	Thr	
225		C1	Δ	1	230 The	A 1	D	C	Λ	235	1	C	Λ	C	240
Lys	Leu	61 y	Asn		Inr	Ala	Pro	Cys		GJU	Lys	Cys	Arg		Ser
Son	Sor	114	Tyr	245 Sor	Son	11.	Cvc	Glu	250	Acr	Acr	116	Glu	255 Typ	Dha
261	961	116	260	oei.	261	116	CyS	265	m g	ush	veb	116	270	1 y 1	1 116
Sor	Pro	Cve		Δla	Gly	Cvc	The		Sor	Lve	Δ1 ₆	Gla	Asn	Gla	Lvc
JU1	110	Uy S	1116	und	O 1 A	ONS	1111	ı yı	oc.i	riz	urg	OHI	USH	0111	- y

Lys Met Tyr Tyr Asn Cys Ser Cys lle Lys Glu Gly Leu lle Thr Ala Asp Ala Glu Gly Asp Phe Ile Asp Ala Arg Pro Gly Lys Cys Asp Ala Lys Cys Tyr Lys Leu Pro Leu Phe Ile Ala Phe Ile Phe Ser Thr Leu Ile Phe Ser Gly Phe Ser Gly Val Pro Ile Val Leu Ala Met Thr Arg Val Val Pro Asp Lys Leu Arg Ser Leu Ala Leu Gly Val Ser Tyr Val lle Leu Arg lle Phe Gly Thr lle Pro Gly Pro Ser lle Phe Lys Met Ser Gly Glu Thr Ser Cys Ile Leu Arg Asp Val Asn Lys Cys Gly His Arg Gly Arg Cys Trp Ile Tyr Asn Lys Thr Lys Met Ala Phe Leu Leu Val Gly Ile Cys Phe Leu Cys Lys Leu Cys Thr Ile Ile Phe Thr Thr Ile Ala Phe Phe Ile Tyr Lys Arg Arg Leu Asn Glu Asn Thr Asp Phe Pro Asp Val Thr Val Lys Asn Pro Lys Val Lys Lys Glu Glu Thr Asp Leu

<210> 4898

<211> 679

<212> PRT

<213> Homo sapiens

<400> 521

Met Met His Tyr Leu Lys Asn Ile Met Ile Ala Val Val Glu Ser Met

1 5 10 15

Ile Asn Lys Phe Glu Glu Asp Glu Thr Arg Asn Gln Glu Arg Gln Lys

			20					25					30		
Lys	Ile	Gln	Lys	Glu	Lys	Ser	His	Ser	Tyr	Arg	Thr	Asp	Asn	Cys	Ser
		35					40					45			
Asp	Ser	Asp	Ser	Ser	Leu	Asn	Gln	Ser	Tyr	Lys	Phe	Cys	Gln	G1 y	Lys
	50					55					60				
Leu	Gln	Leu	lle	Leu	Asp	Gln	Leu	Asp	Pro	Gly	Gln	Pro	Lys	Glu	Val
65					70					75					80
Arg	Tyr	Glu	Ala	Leu	Gln	Thr	Leu	Cys	Ser	Ala	Pro	Pro	Ser	Asp	Val
				85					90					95	
Leu	Asn	Cys	Glu	Asn	Trp	Thr	Thr	Leu	Cys	Glu	Lys	Leu	Thr	Val	Ser
			100					105					110		
Leu	Ser	Asp	Pro	Asp	Pro	Val	Phe	Ser	Asp	Arg	He	Leu	Lys	Phe	Cys
		115					120					125			
Ala	Gln	Thr	Phe	Leu	Leu	Ser	Pro	Leu	His	Met	Thr	Lys	Glu	11e	Tyr
	130					135					140				
Thr	Ser	Leu	Ala	Lys	Tyr	Leu	Glu	Ser	Tyr	Phe	Leu	Ser	Arg	Glu	Asn
145					150					155					160
His	Ile	Pro	Thr	Leu	Ser	Ala	Gly	Val	Asp	Ile	Thr	Asn	Pro	Asn	Met
				165					170					175	
Thr	Arg	Leu	Leu	Lys	Lys	Val	Arg	Leu	Leu	Asn	Glu	Tyr	Gln	Lys	Glu
			180					185					190		
Ala	Pro	Ser	Phe	Trp	lle	Arg	llis	Pro	Glu	Lys	Tyr	Met	Glu	Glu	11e
		195					200					205			
Val	Glu	Ser	Thr	Leu	Ser	Leu	Leu	Thr	Val	Lys	His	Asn	Gln	Ser	His
	210					215					220				
Val	Val	Ser	Gln	Lys	lle	Leu	Asp	Pro	lle	Tyr	Phe	Phe	Ala	Leu	Val
225					230					235					240
Asp	Thr	Lys	Ala	Val	Trp	Phe	Lys	Lys	Trp	Met	His	Ala	His	Tyr	Ser
				245					250					255	
Arg	Thr	Thr	Val	Leu	Arg	Leu	Leu	Glu	Lys	Lys	Tyr	Lys	Ser	Leu	Val
			260					265					270		
Thr	Thr	Ala	He	Gln	Gln	Cys	Val	Gln	Tyr	Phe	Glu	Met	Cys	Lys	Thr
		275					280					285			
Arg	Lys	Ala	Asp	Glu	Thr	Leu	Gl y	His	Ser	Lys	His	Cys	Arg	Asn	Lys
	290					295					300				
Gln	Lys	Thr	Phe	Tyr	Tyr	Leu	Gly	Gln	Glu	Leu	Gln	Tyr	He	Tyr	Phe

305					310					315					320
Пe	His	Ser	Leu	Cys	Leu	Leu	Gly	Arg	Leu	Leu	11e	Tyr	Lys	Gln	G]y
				325					330					335	
Arg	Lys	Leu	Phe	Pro	He	Lys	Leu	Lys	Asn	Lys	Lys	Gly	Leu	Val	Ser
			340					345					350		
Leu	He	Asp	Leu	Leu	Val	Leu	Phe	Thr	Gln	Leu	He	Tyr	Tyr	Ser	Pro
		355					360					365			
Ser	Cys	Pro	Lys	Met	Thr	Ser	Ala	Ala	His	Ser	Glu	Asn	Tyr	Ser	Pro
	370					375					380				
Ala	Ser	Met	Val	Thr	Glu	Val	Leu	Trp	He	Leu	Ser	Asp	Gln	Lys	Glu
385					390					395					400
	Ala	Val	Glu	Cys	Leu	Tyr	Asn	Asn	lle	Val	Пе	Glu	Thr	Leu	Leu
				405					410					415	
Gln	Pro	He	His	Asn	Leu	Met	Lys	Gly	Asn	Glu	Ala	Ser	Pro	Asn	Cys
			420					425					430		
Ser	Glu	Thr	Ala	Leu	Ile	His	Ile	Ala	Gly	lle	Leu	Ala	Arg	Пе	Ala
		435					440					445			
Ser	Val	Glu	Glu	Gly	Leu	He	Leu	Leu	Leu	Tyr	Gly	Ala	Asn	Met	Asn
	450					455			٠		460				
Ser	Ser	Glu	Glu	Ser	Pro	Thr	Gly	Ala	His	lle	He	Ala	Gln	Phe	Ser
465					470					475					480
Lys	Lys	Leu	Leu	Asp	Glu	Asp	lle	Ser	He	Phe	Ser	Gly	Ser	Glu	Met
				485					490					495	
Leu	Pro	Va]	Val	Lys	Gly	Ala	Phe	lle	Ser	Val	Cys	Arg	His	He	Tyr
			500					505					510		
Ser	Thr	Cys	Glu	Gly	Leu	Gln	Val	Leu	He	Thr	Tyr	Asn	Leu	His	Glu
		515					520					525			
Ser	He	Ala	Lys	Ala	Trp	Lys	Lys	Thr	Ser	Leu	Leu	Ser	Glu	Arg	11e
	530					535					540				
Pro	Thr	Pro	Val	Glu	Gly	Ser	Asp	Ser	Val	Ser	Ser	Val	Ser	Gln	Glu
545					550					555					560
Ser	Gln	Asn	11e	Met	Ala	Trp	Glu	Asp	Asn	Leu	Leu	Asp	Asp	Leu	Leu
				565					570					575	
His	Phe	Ala	Ala	Thr	Pro	Lys	Gly	Leu	Leu	Leu	Leu	Gln	Arg	Thr	Gly
			580					585					590		
Δla	He	Asn	Glu	Cvs	Val	Thr	Phe	He	Phe	Asn	Arg	Tvr	Ala	Lvs	Lvs

Leu Gln Val Ser Arg His Lys Lys Phe Gly Tyr Gly Val Leu Val Thr Arg Val Ala Ser Thr Ala Ala Gly Gly Ile Ala Leu Lys Lys Ser Gly Phe Ile Asn Glu Leu Ile Thr Glu Leu Trp Ser Asn Leu Glu Tyr Gly Arg Asp Asp Val Arg Val Thr His Pro Arg Thr Thr Pro Val Asp Pro lle Asp Arg Ser Cys Gln Lys <210> 4899 <211> 1148 <212> PRT <213> Homo sapiens <400> 522 Met Val Leu Asn Leu Tyr Gln Leu Asn Gln Leu Asp Cys Pro Gly Gly Arg Leu lle Gly Gly Trp Glu Asp Asn Pro Phe Lys Gly Asp Leu Lys lle Val Leu Arg Gly Asn His Thr Thr Gln Asp Trp Ala Leu Pro Glu Gly Pro Asn Gln Gly Ala Lys Val Leu Glv Val Phe Gly Glu Leu Asp Leu His Gly Ile Pro His Ser Ile Tyr Lys Thr Lys Leu Ser Glu Thr Ala Phe Ala Gly Ser Lys Val Leu Ser Leu Met Asp Ala Val Asp Trp Gln Glu Gly Glu Glu Ile Val Ile Thr Thr Thr Ser Tyr Asp Phe His Gln Thr Glu Thr Arg Ser Ile Val Lys Ile Leu His Asp His Lys Ile

Leu	He	Leu	Asn	Asp	Ser	Leu	Ser	Tyr	Thr	His	Phe	Ala	Glu	Lys	Tyr
	130					135					140				
His	Val	Pro	Gly	Thr	Gly	Glu	Ser	Tyr	Thr	Leu	Ala	Ala	Asp	Val	Gly
145					150					155					160
lle	Leu	Ser	Arg	Asn	lle	Lys	He	Val	Gly	Glu	Asp	Tyr	Pro	Gly	Trp
				165					170					175	
Ser	Glu	Asp	Ser	Phe	Gly	Ala	Arg	Val	Leu	Val	Gly	Ser	Phe	Thr	Glu
			180					185					190		
Asn	Met	Met	Thr	Phe	Lys	Gly	Asn	Ala	Arg	Ile	Ser	Asn	Val	Glu	Phe
		195					200					205			
Tyr	His	Ser	Gly	G1n	Glu	Gly	Phe	Arg	Asp	Ser	Thr	Asp	Pro	Arg	Tyr
	210					215					220				
Ala	Val	Thr	Phe	Leu	Asn	Leu	Gly	Gln	He	Gln	Glu	His	Gly	Ser	Ser
225					230					235					240
Tyr	He	Arg	Gly	Cys	Ala	Phe	His	His	Gly	Phe	Ser	Pro	Ala	He	Gly
				245					250					255	
Val	Phe	Gly	Thr	Λsp	Gly	Leu	Asp	Ile	Asp	Asp	Asn	Пe	He	His	Phe
			260					265					270		
Thr	Val	Gly	Glu	Gly	Ile	Arg	He	Trp	Gly	Asn	Ala	Asn	Arg	Val	Arg
		275					280					285			
Gly	Asn	Leu	lle	Ala	Leu	Ser	Val	Trp	Pro	Gly	Thr	Tyr	Gln	Asn	Arg
	290					295					300				
Lys	Asp	Leu	Ser	Ser	Thr	Leu	Trp	His	Ala	Ala	He	Glu	He	Asn	Arg
305					310					315					320
Gly	Thr	Asn	Thr	Val	Leu	Gln	Asn	Asn	Val	Val	Ala	Gly	Phe	Gly	Arg
				325					330					335	
Ala	Gly	Tyr	Arg	lle	Asp	Gly	Glu		Cys	Pro	Gly	Gln	Phe	Asn	Pro
			340					345					350		
Va]	G] u	Lys	Trp	Phe	Asp	Asn	Glu	Ala	His	Gly	Gly		Tyr	G1 y	Пe
		355					360					365			
Tyr		Asn	Gln	Asp	Gly	Leu	Pro	G1 y	Cys	Ser	Leu	lle	Gln	Gly	Phe
	370					375					380				
Thr	He	Trp	Thr	Cys		Asp	Tyr	Gly	lle	Tyr	Phe	Gln	Thr	Thr	
385					390					395					400
Ser	Val	His	He		Asn	Val	Thr	Leu		Asp	Asn	Gly	Met		He
				405					410					415	

Phe	Pro	Met	He	Tyr	Met	Pro	Ala	Ala	Ιle	Ser	His	Lys	Ile	Ser	Ser
			420					425					430		
Lys	Asn	Val	Gln	lle	Lys	Ser	Ser	Leu	He	Val	Gly	Ser	Ser	Pro	Gly
		435					440					445			
Phe	Asn	Cys	Ser	Asp	Val	Leu	Thr	Asn	Asp	Asp	Pro	Asn	He	Glu	Leu
	450					455			•		460				
Thr	Ala	Ala	His	Arg	Ser	Pro	Arg	Ser	Pro	Ser	Gly	Gly	Arg	Ser	Gly
465					470					475					480
Ile	Arg	Trp	Pro	Thr	Phe	Ala	Ser	Ala	His	Asn	Met	Ala	Pro	Arg	Lys
				485					490					495	
Pro	His	Ala	Gly	He	Met	Ser	Tyr	Asn	Ala	He	Ser	Gly	Leu	Leu	Asp
			500					505					510		
He	Ser	Gly	Ser	Thr	Phe	Val	Gly	Phe	Lys	Asn	Val	Cys	Ser	Gly	Glu
		515					520					525			
Thr	Asn	Val	lle	Phe	lle	Thr	Asn	Pro	Leu	Asn	Glu	Asp	Leu	Gln	His
	530					535					540				
Pro	lle	His	Val	Lys	Asn	Ile	Lys	Leu	Val	Asp	Thr	Thr	Glu	Gln	Ser
545					550					555					560
Lys	He	Phe	Ile	His	Arg	Pro	Asp	He	Ser	Lys	Val	Asn	Pro	Ser	Asp
				565					570					575	
Cys	Val	Asp	Met	Val	Cys	Asp	Ala	Lys	Arg	Lys	Ser	Phe	Leu	Arg	Asp
			580					585					590		
lle	Лsp	Gly	Ser	Phe	Leu	G1 y	Asn	Ala	Gly	Ser	Val		Pro	Gln	Ala
		595					600					605			
Glu		Glu	Trp	Asp	Gly		Ser	Gln	Val	Gly		Gly	Asp	Tyr	Arg
	610					615					620				
	P.ro	Lys	Ala	Met	Leu	Thr	Phe	Leu	Asn		Ser	Arg	He	Pro	
625					630					635				_	640
Thr	Glu	Lys	Ala		His	Lys	GIy	lle		Arg	Asp	Ser	Thr		Lys
-				645	0.1	•			650	ъ.			0.1	655	
Tyr	Leu	Pro		Trp	Gln	Ser	Tyr		Cys	Phe	Gly	Met		Tyr	Ala
			660					665					670		_
Met	Met		He	GJu	Ser	Leu		Pro	Asp	Thr	Glu		Arg	Arg	Leu
	10	675		7.1		0.1	680	a :	T	,, .		685	7.3		6.3
Ser		Val	Ala	He	Met		Asn	Gly	Tyr	Val		Leu	He	Asn	Gly
	690					695					700				

Pro Gln Asp His Gly Trp Cys Ala Gly Tyr Thr Cys Gln Arg Arg Leu Ser Leu Phe His Ser Ile Val Ala Leu Asn Lys Ser Tyr Glu Val Tyr Phe Thr Gly Thr Ser Pro Gln Asn Leu Arg Leu Met Leu Leu Asn Val Asp His Asn Lys Ala Val Leu Val Gly 11e Phe Phe Ser Thr Leu Gln Arg Leu Asp Val Tyr Val Asn Asn Leu Leu Val Cys Pro Lys Thr Thr lle Trp Asn Ala Gln Gln Lys His Cys Glu Leu Asn Asn His Leu Tyr Lys Asp Gln Phe Leu Pro Asn Leu Asp Ser Thr Val Leu Gly Glu Asn Tyr Phe Asp Gly Thr Tyr Gln Met Leu Tyr Leu Leu Val Lys Gly Thr Ile Pro Val Glu Ile His Thr Ala Thr Val Ile Phe Val Ser Phe Gln Leu Ser Val Ala Thr Glu Asp Asp Phe Tyr Thr Ser His Asn Leu Val Lys Asn Leu Ala Leu Phe Leu Lys Ile Pro Ser Asp Lys Ile Arg Ile Ser Lys Ile Arg Gly Lys Ser Leu Arg Arg Lys Arg Ser Met Gly Phe lle lle Glu lle Glu lle Gly Asp Pro Pro lle Gln Phe lle Ser Asn Gly Thr Thr Gly Gln Met Gln Leu Ser Glu Leu Gln Glu 11e Ala Gly Ser Leu Gly Gln Ala Val Ile Leu Gly Asn Ile Ser Ser Ile Leu Gly Phe Asn Ile Ser Ser Met Ser Ile Thr Asn Pro Leu Pro Ser Pro Ser Asp Ser Gly Trp Ile Lys Val Thr Ala Gln Pro Val Glu Arg Ser Ala Phe Pro Val His His Val Ala Phe Val Ser Ser Leu Leu Val Ile Thr

Gln Pro Val Ala Ala Gln Pro Gly Gln Pro Phe Pro Gln Gln Pro Ser 1000 1005 Val Lys Ala Thr Asp Ser Asp Gly Asn Cys Val Ser Val Gly 11e Thr 1015 1020 Ala Leu Thr Leu Arg Ala Ile Leu Lys Asp Ser Asn Asn Asn Gln Val 1030 1035 1025 1040 Asn Gly Leu Ser Gly Asn Thr Thr Ile Pro Phe Ser Ser Cys Trp Ala 1050 1045 Asn Tyr Thr Asp Leu Thr Pro Leu Arg Thr Gly Lys Asn Tyr Lys Ile 1070 1060 1065 Glu Phe Ile Leu Asp Asn Val Val Gly Val Glu Ser Arg Thr Phe Ser 1080 1085 Leu Leu Ala Glu Ser Val Ser Ser Ser Gly Ser Ser Ser Ser Asn 1095 Ser Lys Ala Ser Thr Val Gly Thr Tyr Ala Gln lle Met Thr Val Val 1110 1115 Ile Ser Cys Leu Val Gly Arg Met Trp Leu Leu Glu Ile Phe Met Ala 1125 1130 1135 Ala Val Ser Thr Leu Asn lle Thr Leu Arg Ser Tyr 1145 1140

<210> 4900

<211> 300

<212> PRT

<213> Homo sapiens

<400> 523

Met His Ser Leu Lys Lys Val Thr Phe Glu Asp Val Ala Ile Asp Phe 1 5 10 15 Thr Gln Glu Glu Trp Ala Met Met Asp Thr Ser Lys Arg Lys Leu Tyr 20 25 30

Arg Asp Val Met Leu Glu Asn Ile Ser His Leu Val Ser Leu Gly Tyr
35 40 45
Gln Ile Ser Lys Ser Tyr Ile Ile Leu Gln Leu Glu Gln Gly Lys Glu

	50					55					60				
Leu	Trp	Arg	Glu	Gly	Arg	Glu	Phe	Leu	Gln	Asp	Gln	Asn	Pro	Asp	Arg
65					70					75					80
Glu	Ser	Ala	Leu	Lys	Lys	Lys	His	Met	lle	Ser	Met	His	Pro	He	Thr
				85					90					95	
Arg	Lys	Asp	Ala	Ser	Thr	Ser	Met	Thr	Met	Glu	Asn	Ser	Leu	He	Leu
			100					105					110		
Glu	Asp	Pro	Phe	Glu	Cys	Asn	Asp	Ser	Gly	Glu	Asp	Cys	Thr	His	Ser
		115					120					125			
Ser	Thr	lle	Thr	Gln	Arg	Leu	Leu	Thr	His	Ser	Gly	Lys	Lys	Pro	Tyr
	130					135					140				
Val	Ser	Lys	Gln	Cys	Gly	Lys	Ser	Leu	Arg	Asn	Leu	Phe	Ser	Pro	Lys
145					150					155					160
Pro	His	Lys	Gln	lle	His	Thr	Lys	Gly	Lys	Ser	Tyr	Gln	Cys	Asn	Leu
				165					170					175	
Cys	Glu	Lys	Ala	Tyr	Thr	Asn	Cys	Phe	Arg	Leu	Arg	Arg	His	Lys	Met
			180					185					190		
Thr	His	Thr	Gly	Glu	Arg	Pro	Tyr	Ala	Cys	His	Leu	Cys	Gly	Lys	Ala
		195					200					205			
Phe	Thr	Gln	Cys	Ser	His	Leu	Arg	Arg	His	Glu	Lys	Thr	His	Thr	Gly
	210					215					220				
Glu	Arg	Pro	Tyr	Lys	Cys	His	Gln	Cys	Gly	Lys	Ala	Phe	Ile	Gln	Ser
225					230					235					240
Phe	Asn	Leu	Arg	Arg	His	Glu	Arg	Thr	His	Leu	G1 y	Lys	Lys	Cys	Tyr
				245					250					255	
Glu	Cys	Asp	Lys	Ser	Gly	Lys	Ala	Phe	Ser	Gln	Ser	Ser	Gly	Phe	Arg
			260					265					270		
Gly	Asn	Lys	lle	lle	His	Thr	Gly	Glu	Lys	Pro	His	Ala	Cys	Leu	Leu
		275					280					285			
Cys		Lys	Ala	Phe	Ser	Leu	Ser	Ser	Asp	Leu					
	290					295					300				

<210> 4901

<211> 1216

<212> PRT

<213> Homo sapiens

<400)> 52	24													
Met	Ser	Gln	Ala	Gly	Asp	Val	Glu	Gly	Pro	Ser	Thr	Gly	Asp	Pro	Val
ł				5					10					15	
Leu	Ser	Pro	Gln	His	Asn	Cys	Glu	Leu	Leu	Gln	Asn	Met	Glu	Gly	Ala
			20					25					30		
Ser	Ser	Met	Pro	Gly	Leu	Ser	Pro	Лsp	Gly	Pro	Gly	Ala	Ser	Ser	Gly
		35					40					45			
Pro	Gly	Val	Arg	Ala	Gly	Ser	Arg	Arg	Lys	lle	Pro	Arg	Lys	Glu	Ala
	50					55					60				
Leu	Arg	Gly	Gly	Ser	Ser	Arg	Ala	Ala	G1 y	Ala	Ala	Glu	Val	Arg	Pro
65					70					75					80
Gly	Val	Leu	Glu	Leu	Leu	Ala	Val	Val	Gln	Ser	Arg	Gly	Ser	Met	Leu
				85					90					95	
Ala	Pro	Gly	Leu	His	Met	Gln	Leu	Pro	Ser	Val	Pro	Thr	Gln	Gly	Arg
			100					105					110		
Ala	Leu	Thr	Ser	Lys	Arg	Leu	Gln	Val	Ser	Leu	Cys	Asp	Ile	Leu	Asp
		115					120					125			
Asp	Ser	Cys	Pro	Arg	Lys	Leu	Cys	Ser	Arg	Ser	Ala	Gly	Leu	Pro	Glu
	130					135					140				
Arg	Ala	Leu	Ala	Cys	Arg	Glu	Arg	Leu	Ala	Gly	Val	Glu	Glu	Val	Ser
145					150					155					160
Cys	Leu	Arg	Pro	Arg	Glu	Ala	Arg	Asp	Gly	Gly	Met	Ser	Ser	Pro	Gly
				165					170					175	
Cys	Asp	Arg	Arg	Ser	Pro	Thr	Leu	Ser	Lys	Glu	Glu	Pro	Pro	G1y	Arg
			180					185					190		
Pro	Leu	Thr	Ser	Ser	Pro	Asp	Pro	Val	Pro	Val	Arg	Val	Arg	Lys	Lys
		195					200					205			
Trp	Arg	Λrg	Gln	Gly	Ala	His	Ser	Glu	Cys	Glu	Glu	Gly	Ala	Gly	Asp
	210					215					220				
Phe	Leu	Trp	Leu	Asp	Gln	Ser	Pro	Arg	Gly	Asp	Asn	Leu	Leu	Ser	Val
225					230					235					240
Gly	Asp	Pro	Pro		Val	Ala	Asp	Leu		Ser	Leu	Gly	Gly	Pro	Cys
				245					250					255	
Arg	Pro	Pro	Ser	Pro	Lys	Asp	Thr	Gly	Ser	Gly	Pro	Gly	Glu	Pro	Gly

			260					265					270		
Gly	Ser	Gly	Ala	Gly	Cys	Ala	Ser	Gly	Thr	Glu	Lys	Phe	Gly	Tyr	Leu
		275					280					285			
Pro	Ala	Thr	Gly	Asp	Gly	Pro	Gln	Pro	Gly	Ser	Pro	Cys	Gly	Pro	Val
	290					295					300				
Gly	Phe	Pro	Val	Pro	Ser	Gly	Gly	Glu	Ser	Leu	Ser	Ser	Ala	Ala	Gln
305					310					315					320
Ala	Pro	Pro	Gln	Ser	Ala	Ala	Leu	Cys	Leu	Gly	Ala	Ser	Ala	Gln	Ala
				325					330					335	
Ser	Ala	Glu	Gln	Gln	Glu	Ala	Val	Cys	Val	Val	Arg	Thr	Gly	Ser	Asp
			340					345					350		
Glu	Gly	Gln	Ala	Pro	Ala	Gln	Asp	Gln	Glu	Glu	Leu	Glu	Ala	Lys	Ala
		355					360					365			
Gln	Pro	Ala	Ser	Arg	Gly	Arg	Leu	Glu	Gln	Gly	Leu	Ala	Ala	Pro	Ala
	370					375					380				
Asp	Thr	Cys	Ala	Ser	Ser	Arg	Glu	Pro	Leu	Gly	Gly	Leu	Ser	Ser	Ser
385					390					395					400
Leu	Asp	Thr	Glu	Ala	Ser	Arg	Ala	Cys	Ser	Gly	Pro	Phe	Met	Glu	Gln
				405					410					415	
Arg	Arg	Ser	Lys	Gly	Thr	Lys	Asn	Leu	Lys	Lys	Gly	Pro	Val	Pro	Cys
			420					425					430		
Ala	Gln	Asp	Arg	Gly	Thr	Asp	Arg	Ser	Ser	Asp	Asn	Ser	His	Gln	Asp
		435					440					445			
Arg	Pro	Glu	Glu	Pro	Ser	Pro	Gly	Gly	Cys	Pro	Arg	Leu	Glu	Glu	Val
	450					455					460				
Lys	lle	Pro	His	Gly	Val	Lys	Leu	Val	Cys	Tyr	Leu	Gly	Ser	Gly	Pro
465					470					475					480
Val	lle	Gln	Leu	Leu	Gly	Ala	lle	Ser	His	Gly	Gln	Ala	Gly	Gly	Gln
				485					490					495	
Leu	Pro	Pro		Leu	Glu	Val	Leu	Glu	Asp	Leu	Met	Glu	Val	Ser	Ser
			500					505					510		
Pro	Ser		Ala	Gln	Arg	Leu		Arg	Lys	Lys	Arg		Met	Va]	Gln
		515					520					525			
G] y		Ala	Gly	Cys	Gln		Phe	Gln	Pro	Ser		Ser	Gly	G1 y	Thr
	530					535					540				
A 3	(1)	Α	13	C 1	C1	I	C'	Λ	D	D)	T	D	Pro	Λ	C*

545					550					555					560
Gly	Ser	Leu	Ala	Leu 565	Gly	Asp	P.ro	Ser	Ser 570	Asp	Pro	Λla	Cys	Ser 575	G1n
Ser	Gly	Pro	Met 580	Glu	Ala	Glu	Glu	Лsp 585	Ser	Leu	Pro	Glu	Gln 590	Pro	Glu
Asp	Ser	Ala 595	Gln	Leu	Gln	Gln	Glu 600	Lys	Pro	Ser	Leu	Tyr 605	lle	Gly	Va]
Arg	Gly 610	Thr	Val	Val	Arg	Ser 615	Met	Gln	Glu	Val	Leu 620	Trp	Thr	Arg	Leu
Arg 625	Glu	Leu	Pro	Asp	Pro 630	Val	Leu	Ser	Glu	Glu 635	Val	Val	Glu	Gly	Ile 640
Ala	Ala	Gly	He	Glu 645	Ala	Ala	Leu	Trp	Asp 650	Leu	Thr	G1n	Gly	Thr 655	Asn
Gly	Arg	Tyr	Lys 660	Thr	Lys	Tyr	Arg	Ser 665	Leu	Leu	Phe	Asn	Leu 670	Arg	Asp
Pro	Arg	Λsn 675	Leu	Asp	Leu	Phe	Leu 680	Lys	Val	Val	His	Gly 685	Asp	Val	Thr
Pro	Tyr 690	Asp	Leu	Val	Arg	Met 695	Ser	Ser	Met	Gln	Leu 700	Ala	Pro	Gln	Glu
Leu 705	Ala	Arg	Trp	Arg	Asp 710	Gln	Glu	Glu	Lys	Arg 715	Gly	Leu	Asn	lle	11e 720
Glu	Gln	Gln	Gln	Lys 725	Glu	Pro	Cys	Arg	Leu 730	Pro	Ala	Ser	Lys	Met 735	Thr
llis	Lys	Gly	Glu 740	Val	Glu	He	Gln	Arg 745	Asp	Met	Asp	Gln	Thr 750	Leu	Thr
Leu	Glu	Asp 755	Leu	Val	Gly	Pro	Gln 760	Met	Phe	Met	Asp	Cys 765	Ser	Pro	Gln
Ala	Leu 770	Pro	He	Ala	Ser	Glu 775	Asp	Thr	Thr	Gly	G1n 780	His	Asp	His	His
Phe 785	Leu	Asp	Pro	Asn	Cys 790	His	He	Cys	Lys	Asp 795	Trp	Glu	Pro	Ser	Asn 800
Glu	Leu	Leu	Gly	Ser 805	Phe	Glu	Ala	Ala	Lys 810	Ser	Cys	Gly	Asp	Asn 815	lle
Phe	Gln	Lys	Ala 820	Leu	Ser	Gln	Thr	Pro 825	Met	Pro	Ala	Pro	G1u 830	Met	Pro
Lvs	Thr	Arg	Glu	Leu	Ser	Pro	Thr	Glu	Pro	Gln	Asp	Arg	Val	Pro	Pro

		835					840					845			
Ser	Gly	Leu	His	Val	Pro	Λla	Ala	Pro	Thr	Lys	Ala	Leu	Pro	Cys	Leu
	850					855					860			•	
Pro	Pro	Trp	Glu	Gly	Val	Leu	Asp	Met	Phe	Ser	lle	Lys	Arg	Phe	Arg
865					870					875					880
Лlа	Arg	Ala	Gln	Leu	Val	Ser	Gly	His	Ser	Cys	Arg	Leu	Val	Gln	Ala
				885					890					895	
Leu	Pro	Thr	Val	He	Arg	Ser	Ala	Gly	Cys	Ile	Pro	Ser	Asn	Ile	Val
			900					905					910		
Trp	Asp	Leu	Leu	Ala	Ser	He	Cys	Pro	Ala	Lys	Ala	Lys	Asp	Val	Cys
		915					920					925			
Val	Val	Arg	Leu	Cys	Pro	His	Gly	Ala	Arg	Asp	Thr	Gln	Asn	Cys	Arg
	930					935					940				
Leu	Leu	Tyr	Ser	Tyr	Leu	Asn	Asp	Arg	Gln	Arg	His	Gly	Leu	Ala	Ser
945					950					955					960
Va]	Glu	His	Met	Gly	Met	Val	Leu	Leu	Pro	Leu	Pro	Ala	Phe	Gln	Pro
				965					970					975	
Leu	Pro	Thr	Arg	Leu	Arg	Pro	Leu	Gly	Gly	Pro	Gly	Leu	Trp	Ala	Leu
			980					985					990		
Pro	Val	Ser	Pro	Leu	Leu	Ser	Pro	Gly	Leu	Glu	Val	Thr	His	Ser	Ser
		995					1000					1005			
Leu	Leu	Leu	Ala	Val	Leu	Leu	Pro	Lys	Glu	Gly	Leu	Pro	Asp	Thr	Ala
	1010				0	1015					1020				
Gly	Ser	Ser	Pro	Trp	Leu	Gly	Lys	Val	Gln	Lys	Met	Val	Ser	Phe	Asn
102	ō				1030					1035					1040
Ser	Lys	Val	Glu	Lys	Arg	Tyr	Tyr	Gln	Pro	Asp	Asp	Arg	Arg	Pro	Asn
				1045					1050					1055	
Va]	Pro	Leu	Lys	Gly	Thr	Pro	Pro	Pro	Gly	Gly	Ala	Trp	Gln	Gln	Ser
			1060					1065					1070		
Gln			Gly	Ser	lle			Arg	Gly	lle		Ala	Trp	Gln	Λrg
		1075					1080					1085			
Pro	Pro	Arg	Gly	Arg			Leu	Trp	Pro	Glu	Pro	Glu	Asn	Trp	Gln
	1090					1095					1100				
		Gly	Arg			Trp	Pro	Pro			Gly	Leu	Arg		
110	5				1110					1115					1120
(1	Ulio	Unc	Tur	Car	Mos I	Ala	Dr.c	Alc	(1	III i ~	C1	Dh.	(1	A >= c=	1:1.

Gln His Phe His Arg Asp Ser Cys Pro His Gln Ala Leu Leu Arg His Leu Glu Ser Leu Ala Thr Met Ser His Gln Leu Gln Ala Leu Leu Cys Pro Gln Thr Lys Ser Ser 11e Pro Arg Pro Leu Gln Arg Leu Ser Ser Ala Leu Ala Ala Pro Glu Pro Pro Gly Pro Ala Arg Asp Ser Ser Leu Gly Pro Thr Asp Glu Ala Gly Ser Glu Cys Pro Phe Pro Arg Lys Ala

<210> 4902

<211> 435

<212> PRT

<213> Homo sapiens

<400> 525 Met Arg Lys Leu Leu Thr Asn Leu Pro Ala Ala Ala Val Leu Ser Ala Gln Val Tyr Ser Ala Val Leu Arg Gly Leu Trp Glu Glu Asn Val Cys Gly Thr Pro Gly Arg Thr Arg Val Cys Thr Ala Leu Leu Tyr Gly Gln Val Cys Pro Phe Gln Asp Ser Thr Asp Gly Leu Arg Thr 11e Thr Ser lle Leu Phe Asn Trp Pro Pro Glu Asn Thr Ser Val Tyr Tyr Gln Pro Pro Gln Arg Ser Ser Phe Arg lle Lys Leu Ala Phe Arg Asn Leu Scr Trp Pro Gly Leu Gly Leu Glu Asp His Gln Glu 11e Val Leu Gly Gln Leu Val Leu Pro Glu Pro Asn Glu Ala Lys Pro Asp Asp Pro Ala Pro

Arg Pro Gly Gln His Ala Leu Thr Met Pro Ala Leu Glu Pro Ala Pro

	130					135					140				
Pro	Leu	Leu	Λla	Лѕр	Leu	Gly	Pro	Ala	Leu	Glu	Pro	Glu	Ser	Pro	Ala
145					150					155					160
Ala	Leu	Gly	Pro	Pro	Gly	Tyr	Leu	His	Ser	Ala	Pro	Gly	Pro	Ala	Pro
				165					170					175	
Ala	Pro	Gly	Glu	Glu	Pro	Pro	Pro	Gly	Thr	Val	Leu	Glu	Pro	Gln	Se.r
			180					185					190		
Ala	Pro	Glu	Ser	Ser	Cys	Pro	Cys	Arg	Gly	Ser	Val	Lys	Asn	Gln	Pro
		195					200					205			
Ser	Glu	Glu	Leu	Pro	Asp	Met	Thr	Thr	Phe	Pro	Pro	Arg	Leu	Leu	Ala
	210					215					220				
Glu	Gln	Leu	Thr	Leu	Met	Asp	Ala	Glu	Leu	Phe	Lys	Lys	Va]	Val	Leu
225					230					235					240
Tyr	Glu	Cys	Leu	Gly	Cys	He	Trp	Gly	Gln	G1 y	His	Leu	Lys	Gly	Asn
				245					250					255	
Glu	His	Met	Ala	Pro	Thr	Val	Arg	Ala	Thr	He	Ala	His	Phe	Asn	Arg
			260					265					270		
Leu	Thr	Asn	Cys	Ile	Thr	Thr	Ser	Cys	Leu	Gly	Asp	His	Ser	Met	Arg
		275					280					285			
Ala	Arg	Asp	Arg	Ala	Arg	Va]	Val	Glu	His	Trp	Ile	Lys	Val	Ala	Arg
	290					295					300				
Glu	Cys	Leu	Ser	Leu	Asn	Asn	Phe	Ser	Ser	Va]	His	Val	Ile	Val	Ser
305					310					315					320
Ala	Leu	Cys	Ser	Asn	Pro	He	G1 y	Gln	Leu	llis	Lys	Thr	Trp	Ala	G1 y
				325					330					335	
Val	Ser	Ser	Lys	Ser	Met	Lys	Glu	Leu	Lys	Glu	Leu	Cys	Lys	Lys	Asp
			340										350		
Thr	Ala		Lys	Arg	Asp	Leu		He	Lys	Arg	Asp		Gly	Glu	Asn
		355					360					365			
Asn		Àsn	Ser	Tyr	Arg		Leu	Ser	Pro	Gln		Pro	Cys	Glu	Val
	370					375					380				
	Val	Val	Cys	lle		Leu	Tyr	Arg	Arg		Asn	Ser	Leu	Arg	
385					390					395					400
Ala	Arg	Leu	Gln		Trp	Va]	Thr	His		Gly	Ser	Val	Asp	Leu	Gly
	_			405					410					415	
Gln	Trp	Arg	Gly	Ala	Leu	Arg	Trp	Pro	Pro	Arg	Arg	Gly	Thr	Pro	Arg

Glu Pro Arg <210> 4903 <211> 171 <212> PRT <213> Homo sapiens <400> 526 Met Gly Lys Ser Glu Ile Lys Arg Thr Gly Gly Phe Gly Ser Thr Asn Lys Gln Gly Lys Ala Ala Tyr Trp Val Asn Gln 11e Thr Asp Lys Cys Pro Thr Cys Glu Ile Thr Ile Gln Gly Lys Lys Phe Lys Gly Leu Val Asp Thr Arg Ala Asp Ile Ser Ile Ile Ser Leu Gln His Trp Pro Ser Thr Trp Pro Ile Gln Pro Thr Gln Phe Asn lle Val Gly Val Gly Glu Ala Pro Glu Val Tyr Gln Ser Ser Ser Val Leu Pro Cys Glu Gly Pro Asp Gly Gln Pro Glu Thr He Gln Pro He He Thr Ser Val Ser He Asn Leu Trp Gly Arg Asp Leu Leu Gln Gln Cys Arg Ala Gln Val Leu lle Pro Glu Gln Leu Tyr Ser Pro Gln Ser Gln His Met Met His Glu Met Gly Tyr Val Pro Gly Met Gly Leu Gln Lys Asn Leu Gln Gly Leu Lys Ser Ser Arg Gln Arg Leu Gly Asn Asn Phe

<211> 599

```
<212> PRT
<213> Homo sapiens
<400> 527
Met Asp Pro Pro Met Asp Asp Gln Pro Gly Glu Lys Glu Leu Val Lys
                  5
                                     10
Arg Ser Gln Leu Asp Gly Glu Gly Asp Gly Pro Leu Ser Asn Gln Leu
                                 25
Ser Ala Ser Ser Thr Ile Asn Pro Val Pro Leu Val Gly Leu Gln Lys
         35
                             40
                                                  45
Pro Glu Met Ser Leu Pro Val Lys Pro Gly Gln Gly Asp Ser Glu Ala
                         55
Ser Ser Pro Phe Thr Pro Val Ala Asp Glu Asp Ser Val Val Phe Ser
65
                     70
                                          75
Lys Leu Thr Tyr Leu Gly Cys Ala Ser Val Asn Ala Pro Arg Ser Glu
                                     90
Val Glu Ala Leu Arg Met Met Ser Ile Leu Arg Ser Gln Cys Gln Ile
                                105
                                                     110
Ser Leu Asp Val Thr Leu Ser Val Pro Asn Val Ser Glu Gly Ile Val
        115
                            120
                                                 125
Arg Leu Leu Asp Pro Gln Thr Asn Thr Glu lle Ala Asn Tyr Pro Ile
                        135
Tyr Lys Ile Leu Phe Cys Val Arg Gly His Asp Gly Thr Pro Glu Ser
                    150
                                         155
                                                             160
Asp Cys Phe Ala Phe Thr Glu Ser His Tyr Asn Ala Glu Leu Phe Arg
                165
                                     170
lle His Val Phe Arg Cys Glu lle Gln Glu Ala Val Ser Arg lle Leu
                                185
                                                     190
Tyr Ser Phe Ala Thr Ala Phe Arg Arg Ser Ala Lys Gln Thr Pro Leu
        195
                            200
                                                 205
Ser Ala Thr Ala Ala Pro Gln Thr Pro Asp Ser Asp 11e Phe Thr Phe
                        215
Ser Val Ser Leu Glu Ile Lys Glu Asp Asp Gly Lys Gly Tyr Phe Ser
225
                    230
                                         235
                                                             240
```

Ala Val Pro Lys Asp Lys Asp Arg Gln Cys Phe Lys Leu Arg Gln Gly

				245					250					255	
He	Asp	Lys	Lys	Ile	Val	He	Tyr	Val	Gln	Gln	Thr	Thr	Asn	Lys	Glu
			260					265					270		
Leu	Ala	Пe	Glu	Arg	Cys	Phe	Gly	Leu	Leu	Leu	Ser	Pro	Gly	Lys	Asp
		275					280					285			
Val	Arg	Asn	Ser	Asp	Met	His	Leu	Leu	Asp	Leu	Glu	Ser	Met	Gly	Lys
	290					295					300				
Ser	Ser	Asp	G1y	Lys	Ser	Tyr	Val	lle	Thr	Gly	Ser	Trp	Asn	Pro	Lys
305					310					315					320
Ser	Pro	His	Phe	Gln	Val	Val	Asn	Glu	Glu	Thr	Pro	Lys	Asp	Lys	Val
				325					330					335	
Leu	Phe	Met	Thr	Thr	Ala	Val	Asp	Leu	Val	He	Thr	Glu	Va]	Gln	Glu
			340					345					350		
Pro	Val	Arg	Phe	Leu	Leu	Glu	Thr	Lys	Val	Arg	Val	Cys	Ser	Pro	Asn
		355					360					365			
Glu	Arg	Leu	Phe	Trp	Pro	Phe	Ser	Lys	Arg	Ser	Thr	Thr	Glu	Asn	Phe
•	370					375					380				
Phe	Leu	Lys	Leu	Lys	Gln	Ile	Lys	Gln	Arg	Glu	Arg	Lys	Asn	Asn	Thr
385					390					395					400
Asp	Thr	Leu	Tyr	Glu	Val	Val	Cys	Leu	Glu	Ser	Glu	Ser	Glu	Arg	Glu
				405					410					415	
Arg	Arg	Lys	Thr	Thr	Ala	Ser	Pro	Ser	Val	Arg	Leu	Pro	Gln	Ser	Gly
			420					425					430		
Ser	Gln	Ser	Ser	Val	lle	Pro	Ser	Pro	Pro	Glu	Asp	Asp	Glu	Glu	Glu
		435					440					445			
Asp	Asn	Asp	Glu	Pro	Leu	Leu	Ser	Gly	Ser	Gly	Asp	Val	Ser	Lys	Glu
	450					455					460				
Cys	Ala	Glu	Lys	He	Leu	Glu	Thr	Trp	Gly	Glu	Leu	Leu	Ser	Lys	Trp
465					470					475					480
His	Leu	Asn	Leu	Asn	Val	Arg	Pro	Lys	G1n	Leu	Ser	Ser	Leu	Val	Arg
				485					490					495	
Asn	Gly	Val	Pro	Glu	Ala	Leu	Arg	Gly	Glu	Val	Trp	Gln	Leu	Leu	Ala
			500					505					510		
Gly	Cys	llis	Asn	Asn	Asp	His	Leu	Val	Glu	Lys	Tyr	Arg	lle	Leu	He
		515					520					525			
Thr	Lys	Glu	Ser	Pro	Gln	Asp	Ser	Λla	He	Thr	Arg	Asp	He	Asn	Arg

Thr Phe Pro Ala His Asp Tyr Phe Lys Asp Thr Gly Gly Asp Gly Gln Asp Ser Leu Tyr Lys Ile Cys Lys Val Phe His Val Lys Lys Lys Asp Ser Ile Leu Ser Gly Gly Ser Thr Leu Lys Leu His Lys Lys Gln Leu Gln Ser Val Ile Cys Ile <210> 4905 <211> 1340 <212> PRT <213> Homo sapiens <400> 528 Met Leu Arg Lys Gly Ala Asn Arg Tyr Leu Thr Val Lys Lys Asp Gly Ser Glu Thr Ala His Ala Met Met Thr Cys Asn Leu Thr His Asn Thr Lys His Ala Val Arg Ser Leu Ile Gln Arg Phe Pro Val Thr Asn Lys Glu Arg Thr Glu Leu Leu Pro Lys Thr Glu Arg Gly Asn Val Phe Ala Val Glu Ala Glu Asn Arg Glu Met Ser Lys Thr Ser Gly Arg Leu Asn Asn Gly 11e Pro Gln 11e Pro Val Lys Arg Gly Glu Ser Glu Phe Asp Ser Phe Arg Gln Ser Leu Pro Val Phe Glu Lys Gln Glu Glu Ile Val Lys Ile Ile Lys Glu Asn Lys Val Val Leu Ile Val Gly Glu Thr Gly Ser Gly Lys Thr Thr Gln 11e Pro Gln Phe Leu Leu Asp Asp Cys

Phe Lys Asn Gly Ile Pro Cys Arg Ile Phe Cys Thr Gln Pro Arg Arg

145					150					155					160
Leu	Ala	Ala	lle	Λla	Val	Ala	Glu	Arg	Val	Ala	Ala	Glu	Arg	Arg	Glu
				165					170					175	
Arg	He	Gly	Gln	Thr	He	Gly	Tyr	Gln	He	Arg	Leu	Glu	Ser	Arg	Val
			180					185					190		
Ser	Pro	Lys	Thr	Leu	Leu	Thr	Phe	Cys	Thr	Asn	Gly	Val	Leu	Leu	Arg
		195					200					205			
Thr	Leu	Met	Ala	Gly	Asp	Ser	Thr	Leu	Ser	Thr	Val	Thr	His	Val	He
	210					215					220				
Val	Asp	Glu	Val	His	Glu	Arg	Asp	Arg	Phe	Ser	Asp	Phe	Leu	Leu	Thr
225					230					235					240
Lys	Leu	Arg	Asp	Leu	Leu	Gln	Lys	His	Pro	Thr	Leu	Lys	Leu	He	Leu
				245					250					255	
Ser	Ser	Ala	Ala	Leu	Asp	Val	Asn	Leu	Phe	He	Arg	Tyr	Phe	Gly	Ser
			260					265					270		
Cys	Pro	Val	lle	Tyr	Ile	Gln	Gly	Arg	Pro	Phe	Glu	Val	Lys	Glu	Met
		275					280					285			
Phe	Leu	Glu	Asp	Ile	Leu	Arg	Thr	Thr	Gly	Tyr	Thr	Asn	Lys	Glu	Met
	290					295					300				
Leu	Lys	Tyr	Lys	Lys	Glu	Lys	Gln	Gln	Glu	Glu	Lys	Gln	Gln	Thr	Thr
305					310					315					320
Leu	Thr	Glu	Trp	Tyr	Ser	Ala	Gln	Glu	Asn	Ser	Phe	Lys	Pro	Glu	Ser
				325					330					335	
Gln	Arg	Gln	Arg	Thr	Val	Leu	Asn	Val	Thr	Asp	Glu	Tyr	Asp	Leu	Leu
			340					345					350		
Asp	Asp	Gly	Gly	Лsp	Ala	Va]	Phe	Ser	Gln	Leu	Thr	Glu	Lys	Asp	Val
		355					360					365			
Asn	Cys	Leu	Glu	Pro	Trp	Leu	He	Lys	Glu	Met	Asp	Ala	Cys	Leu	Ser
	370			\		375					380				
Asp	lle	Trp	Leu	His	Lys	Asp	lle	Asp	Ala	Phe	Ala	Gln	Val	Phe	His
385					390					395					400
Leu	He	Leu	Thr	Glu	Asn	Val	Ser	Va]	Asp	Tyr	Arg	His	Ser	Glu	Thr
				405					410					415	
Ser	Ala	Thr	Ala	Leu	Met	Val	Ala	Ala	Gly	Arg	Gly	Phe	Ala	Ser	Gln
			420					425					430		
Val	Glu	Gln	Leu	He	Ser	Met	Gly	Ala	Asn	Val	His	Ser	Lys	Ala	Ser

		435					440					445			
Asn	Gly	Trp	Met	Ala	Leu	Asp	Trp	Ala	Lys	His	Phe	Gly	Gln	Thr	Glu
	450					455					460				
lle	Val	Asp	Leu	Leu	Glu	Ser	Tyr	Ser	Ala	Ser	Leu	Glu	Phe	Gly	Asn
465					470					475					480
Leu	Asp	Glu	Ser	Ser	Leu	Val	Gln	Thr	Asn	Gly	Ser	Лsp	Leu	Ser	Ala
				485					490					495	
Glu	Asp	Arg	Glu	Leu	Leu	Lys	Ala	Tyr	His	His	Ser	Phe	Asp	Asp	Glu
			500					505					510		
Lys	Val	Asp	Leu	Asp	Leu	Ile	Met	His	Leu	Leu	Tyr	Asn	Ile	Cys	His
		515					520					525			
Ser	Cys	Asp	Ala	Gly	Ala	Val	Leu	lle	Phe	Leu	Pro	Gly	Tyr	Asp	Glu
	530					535					540				
He	Val	Gly	Leu	Arg	Asp	Arg	lle	Leu	Phe	Asp	Asp	Lys	Arg	Phe	Ala
545					550					555					560
Asp	Asn	Thr	His	Arg	Tyr	Gln	Val	Phe	Met	Leu	His	Ser	Asn	Met	Gln
				565					570					575	
Thr	Ser	Asp	Gln	Lys	Lys	Val	Leu	Lys	Asn	Pro	Pro	Ala	Gly	Val	Arg
			580					585					590		
Lys	lle	lle	Leu	Ser	Thr	Asn	lle	Ala	Glu	Thr	Ser	11e	Thr	Val	Asn
		595					600					605			
Asp	Val	Val	Phe	Val	Ile	Asp	Ser	Gly	Lys	Val	Lys	Glu	Lys	Ser	Phe
	610					615					620				
Asp	Ala	Leu	Asn	Phe	Va]	Thr	Met	Leu	Lys	Met	Val	Trp	lle	Ser	Lys
625					630					635					640
Ala	Ser	Ala	lle	Gln	Arg	Lys	Gly	Arg	Ala	Gly	Arg	Cys	Arg	Pro	Gly
				645					650					655	
lle	Cys	Phe	Arg	Leu	Phe	Ser	Arg	Leu	Arg	Phe	Gln	Asn	Met	Leu	Glu
			660					665					670		
Phe	Gln	Thr	Pro	Glu	Leu	Leu	Arg	Met	Pro	Leu	Gln	Glu	Leu	Cys	Leu
		675					680					685			
His	Thr	Lys	Leu	Leu	Ala	Pro	Val	Asn	Cys	Pro	He	Ala	Asp	Phe	Leu
	690					695					700				
Met	Lys	Ala	Pro	Glu	Pro	Pro	Pro	Āla	Leu		Val	Arg	Asn	Ala	Val
705					710					715					720
Gln	Met	Leu	Lys	Thr	lle	Asp	Ala	Met	Asp	Thr	Trp	Glu	Asp	Leu	Thr

				725					730					735	
Glu	Leu	Gly	Tyr	His	Leu	Ala	Asp	Leu	Pro	Val	Glu	Pro	His	Leu	Gly
			740					745					750		
Lys	Met	Val	Leu	Cys	Ala	Val	Val	Leu	Lys	Cys	Leu	Asp	Pro	He	Leu
		755					760					765			
Thr	lle	Ala	Cys	Thr	Leu	Ala	Tyr	Arg	Asp	Pro	Phe	Val	Leu	Pro	Thr
	770					775					780				
Gln	Ala	Ser	Gln	Lys	Arg	Ala	Ala	Met	Leu	Cys	Arg	Lys	Arg	Phe	Thr
785					790					795					800
Ala	Gly	Ala	Phe	Ser	Asp	His	Met	Ala	Leu	Leu	Arg	Ala	Phe	Gln	Ala
				805					810					815	
Trp	Gln	Lys	Ala	۸rg	Ser	Asp	Gly	Trp	Glu	Arg	Ala	Phe	Cys	Glu	Lys
			820					825					830		
Asn	Phe	Leu	Ser	Gln	Ala	Thr	Met	Glu	He	He	He	Gly	Met	Arg	Thr
		835					840					845			
Gln	Leu	Leu	Gly	Gln	Leu	Arg	Ala	Ser	Gly	Phe	Val	Arg	Ala	Arg	Gly
	850					855					860				
Gly	Gly	Asp	lle	Arg	Asp	Val	Asn	Thr	Asn	Ser	Glu	Asn	Trp	Ala	Val
865					870					875					880
Val	Lys	Ala	Ala	Leu	Va]	Ala	Gly	Met	Tyr	Pro	Asn	Leu	Va]	His	Val
				885					890					895	
Asp	Arg	Glu	Asn	Leu	Val	Leu	Thr	Gly	Pro	Lys	Glu	Lys	Lys	Val	Arg
			900					905					910		
Phe	His	Pro	Ala	Ser	Val	Leu	Ser	Gln	Pro	Gln	Tyr	Lys	Lys	lle	Pro
		915					920					925			
Pro	Ala	Asn	Gly	Gln	Ala	Ala	Ala	He	Lys	Ala	Leu	Pro	Thr	Asp	Trp
	930					935					940				
Leu	lle	Tyr	Asp	Glu	Met	Thr	Arg	Ala	His	Arg	lle	Ala	Asn	He	Arg
945					950					955					960
Cys	Cys	Ser	Ala	Val	Thr	Pro	Val	Thr		Leu	Val	Phe	Cys	Gly	Pro
				965					970					975	
Ala	Arg	Leu	Ala	Ser	Asn	Ala	Leu	Gln	Glu	Pro	Ser	Ser	Phe	Arg	Val
			980					985					990		
Asp	Gly		Pro	Λsn	Asp			Asp	Ser	Glu		Glu	Asp	Lys	Thr
		995					1000					1005			_
Thr	Ala	Asn	Leu	Ala	Ala	Leu	Lys	Leu	Asp	Glu	Trp	Leu	His	Phe	Thr

1010]	1015]	1020				
Leu Glu	Pro	Glu	Ala	Ala	Ser	Leu	Leu	Leu	Gln	Leu	Arg	Gln	Lys	Trp
1025			1	030]	035				1	040
His Ser	Leu	Phe	Leu	Arg	Arg	Met	۸rg	Ala	Pro	Ser	Lys	Pro	Trp	Ser
]	045				1	050				1	055	
Gln Val	Asp	Glu	Ala	Thr	lle	Arg	Ala	11e	He	Ala	Val	Leu	Ser	Thr
]	060]	065					1070		
Glu Glu	Gln	Ser	Ala	Gly	Leu	Gln	Gln	Pro	Ser	Gly	Ile	Gly	Gln	Arg
1	1075					1080]	085			
Pro Arg	Pro	Met	Ser	Ser	Glu	Glu	Leu	Pro	Leu	Ala	Ser	Ser	Trp	Arg
1090		•		j	1095					1100				
Ser Asn	Asn	Ser	Arg	Lys	Ser	Ser	Ala	Asp	Thr	Glu	Phe	Ser	Asp	Glu
1105]	1110]	1115]	120
Cys Thr	Thr	Ala	Glu	Arg	Val	Leu	Met	Lys	Ser	Pro	Ser	Pro	Ala	Leu
]	1125					1130]	135	
His Pro	Pro	Gln	Lys	Tyr	Lys	Asp	Arg	Gly	Ile	Leu	His	Pro	Lys	Arg
	J	140					1145					1150		
Gly Thr	Glu	Asp	Arg	Ser	Asp	Gln	Ser	Ser	Leu	Lys	Ser	Thr	Asp	Ser
1	1155					1160					1165			
Ser Ser	Tyr	Pro	Ser	Pro	Cys	Ala	Ser	Pro	Ser	Pro	Pro	Ser	Ser	Gly
1170					1175					1180				
Lys Gly	Ser	Lys	Ser	Pro	Ser	Pro	Arg	Pro	Asn	Met	Pro	Val	Arg	Tyr
1185				1190					1195]	1200
Phe lle	Met	Lys	Ser	Ser	Asn	Leu	Arg	Asn	Leu	Glu	He	Ser	Gln	Gln
		1	1205					1210]	1215	
Lys Gly	lle	Trp	Ser	Thr	Thr	Pro	Ser	Asn	Glu	Arg	Lys	Leu	Asn	Arg
]	1220					1225					1230		
Ala Phe	Trp	Glu	Ser	Ser	11e	Va]	Tyr	Leu	Val	Phe	Ser	Val	Gln	Gly
1	1235					1240					1245			
Ser Gly	His	Phe	Gln	Gly	Phe	Ser	Arg	Met	Ser	Ser	Glu	Ile	Gly	Arg
1250					1255					1260				
Glu Lys	Ser	Gln	Asp	Trp	Gly	Ser	Ala	Gly	Leu	Gly	Gly	Val	Phe	Lys
1265				1270					1275				-	1280
Val Glu	Trp	He	Arg	Lys	Glu	Ser	Leu	Pro	Phe	Gln	Phe	Ala	His	His
			1285					1290					1295	

Leu Leu Asn Pro Trp Asn Asp Asn Lys Lys Val Gln Ile Ser Arg Asp
1300
1305
1310

Gly Gln Glu Leu Glu Pro Gln Val Gly Glu Gln Leu Leu Gln Leu Trp
1315
1320
1325

Glu Arg Leu Pro Leu Gly Glu Lys Asn Thr Thr Asp
1330
1335
1340

<210> 4906

<211> 574

<212> PRT

<213> Homo sapiens

<400> 529 Met Thr Ala Gly Ser Val Cys Val Pro Gln lle lle Pro Leu Arg Val 1 5 10 Pro Gln Pro Gly Lys Ala Asn His Glu Ile Asp Asn Asn Thr Leu Leu 25 Glu Met Lys Ser Asp Thr Pro Asp Val Asn Ile Tyr Tyr Thr Leu Asp 45 35 40 Gly Ser Lys Pro Glu Phe Leu Lys Arg lle Gly Tyr Gly Glu Asn Asn 55 Thr Phe Lys Tyr lle Lys Pro lle Thr Leu Pro Asp Gly Lys Ile Gln 70 75 Val Lys Ala lle Ala Val Ser Lys Asp Cys Arg Gln Ser Gly lle Val 85 90 95 Thr Lys Val Phe His Val Asp Tyr Glu Pro Pro Asn Ile Val Ser Pro 105 Glu Asp Asn Val Glu Asn Val Leu Lys Asp Ser Ser Arg Gln Glu Phe 115 125 120 Lys Asn Gly Phe Val Gly Ser Lys Leu Lys Lys Lys Tyr Lys Asn Ser 130 135 Glu Asn Gln Arg Ser Trp Asn Val Asn Leu Arg Lys Phe Pro Glu Ser

Pro Leu Glu lle Pro Ala Tyr Gly Gly Gly Ser Gly Ser Arg Pro Pro

155

175

170

150

Thr	Arg	Gln		Gln	Ser	Pro	Gly		Ala	His	Val	Ser		Gln	Lys
			180					185					190		
Cys	Leu	Thr	Ser	Thr	Glu	He	Met	Arg	He	Gln	Arg	Glu	Thr	Asp	Phe
		195					200					205			
Leu	Lys	Cys	Ala	llis	Cys	Leu	Ala	Pro	Arg	Pro	Ser	Asp	Pro	Phe	Ala
	210					215					220				
Arg	Phe	Cys	Gln	Glu	Cys	Gly	Ser	Pro	Val	Pro	Pro	lle	Phe	Gly	Cys
225					230					235					240
Arg	Leu	Pro	Pro	Pro	Glu	Gly	Ala	Gln	Met	Gly	Leu	Cys	Ala	Glu	Cys
				245					250					255	
Arg	Ser	Leu	Val	Pro	Met	Asn	Thr	Pro	lle	Cys	Val	Val	Cys	Glu	Ala
			260					265					270		
Pro	Leu	Ala	Leu	Gln	Leu	Gln	Pro	Gln	Λla	Ser	Leu	His	Leu	Lys	Glu
		275					280					285			
Lys	Val	lle	Cys	Arg	Ala	Cys	Gly	Thr	Gly	Asn	Pro	Ala	His	Leu	Arg
	290					295					300				
Tyr	Cys	Val	Thr	Cys	Glu	Gly	Ala	Leu	Pro	Ser	Ser	Gln	Glu	Ser	Met
305					310					315					320
Cys	Ser	Gly	Asp	Lys	Ala	Pro	Pro	Pro	Pro	Thr	Gln	Lys	Gly	Gly	Thr
				325					330					335	
Пе	Ser	Cys	Tyr	Arg	Cys	Gly	Arg	Trp	Asn	Leu	Trp	Glu	Ala	Ser	Phe
			340					345					350		
Cys	Gly	Trp	Cys	Gly	Ala	Met	Leu	Gly	11e	Pro	Ala	Gly	Cys	Ser	Val
		355					360					365			
Cys	Pro	Lys	Cys	Gly	Ala	Ser	Asn	His	Leu	Ser	Ala	Arg	Phe	Cys	Gly
	370					375					380				
Ser	Cys	Gly	He	Cys	Val	Lys	Ser	Leu	Val	Lys	Leu	Ser	Leu	Asp	Arg
385					390					395					400
Ser	Leu	Ala	Leu	Ala	Ala	Glu	Glu	Pro	Arg	Pro	Phe	Ser	Glu	Ser	Leu
				405					410					415	
Asn	Пе	Pro	Leu	Pro	Arg	Ser	Asp	Val	Gly	Thr	Lys	Arg	Asp	lle	Gly
			420					425					430		
Thr	Gln	Thr	Va]	Gly	Leu	Phe	Tyr	Pro	Ser	Gly	Lys	Leu	Leu	Ala	Lys
		435					440					445			
Lys	Glu	Gln	Glu	Leu	Ala	Ser	Gln	Lys	Gln	Arg	Gln	Glu	Lys	Met	Ser
	450					455				-	460				

Asp His Lys Pro Leu Leu Thr Ala Ile Ser Pro Gly Arg Gly Tyr Trp 470 475 480 Arg Arg Gln Leu Asp His Ile Ser Ala His Leu Arg Cys Tyr Ala Gln 485 490 Asn Asn Pro Glu Phe Arg Ala Leu Ile Ala Glu Pro Arg Met Gly Lys 500 505 510 Leu lle Ser Ala Thr Val His Glu Asp Gly Cys Glu Val Ser Ile Arg 520 525 Leu Asn Tyr Ser Gln Val Ser Asn Lys Val Arg Lys Leu Arg Leu Arg 530 535 540 Glu Val Lys Gln Pro Ala Ser Ser Lys Gly Thr Lys Leu Val Ser Gly 550 555 560 Arg Pro Arg lle His Thr Trp Gln Pro Glu Thr Phe Pro Ser 570 565

<210> 4907

<211> 534

<212> PRT

<213> Homo sapiens

<400> 530

Met Gly Val Pro Thr Ala Val Ser Ala Thr Pro Val Arg Ala Asp Ala

1 5 10 15

Ser Ser Lys Pro Gln Pro Leu Leu Gln Ser Gln Pro His Leu Phe Phe 20 25 30

Phe Pro Lys Leu Leu Ser Arg Leu Leu Gly Ser Pro Leu Pro Val His
35 40 45

Ser Ala Gly Pro Gly Pro Leu Leu Thr Arg Met Pro Gln Ala Thr Thr 50 55 60

Val Ser Leu Arg Leu Gly Ser Trp Ser Leu Thr Glu Asp Arg Asp Val
65 70 75 80

Ser Gly Glu Trp Pro Arg Ala Phe Pro Asp Thr Pro Pro Gly Met Thr
85 90 95

Thr Ser Val Phe Pro Val Ala Asp Ala Cys His Ser Val Lys Ser Leu 100 105 110

Gln	Arg	Gln	Pro	Gly	Ala	Ser	Pro	Ser	Gln	Glu	Arg	Lys	Pro	Thr	Gly
		115					120					125			
Val	Ser	Val	He	Tyr	Trp	Glu	Arg	Leu	Leu	Leu	Gly	Ser	Arg	Ser	Λsp
	130					135					140				
Gln	Ala	Ser	Пе	Ser	Leu	Arg	Leu	Thr	Ser	Pro	Leu	Arg	Pro	Pro	Lys
145					150					155					160
Ser	Ser	Arg	Pro	Arg	Glu	Lys	Thr	Phe	Thr	Glu	Tyr	Arg	Val	Pro	Gly
				165					170					175	
Arg	Gln	Pro	Arg	Thr	Pro	Glu	Arg	Gln	Lys	Pro	Cys	Ala	Gln	Glu	Val
			180					185					190		
Pro	Gly	Arg	Ala	Phe	Gly	Asn	Ala	Ser	Asp	Leu	Lys	Ala	Ala	Ser	Gly
		195					200					205			
Gly	Arg	Asp	Arg	Arg	Met	Gly	Ala	Лlа	Trp	Gln	Glu	Pro	His	Arg	Leu
	210					215					220				
Leu	Gly	Gly	Gln	Glu	Pro	Ser	Thr	Trp	Asp	Glu	Leu	Gly	Glu	Ala	Leu
225					230					235					240
His	Ala	Gly	Glu	Lys	Ser	Phe	Glu	Cys	Arg	Ala	Cys	Ser	Lys	Val	Phe
				245					250					255	
Val	Lys	Ser	Ser	Asp	Leu	Leu	Lys	His	Leu	Arg	Thr	His	Thr	Gly	Glu
			260					265					270		
Arg	Pro	Tyr	Glu	Cys	Thr	Gln	Cys	Gly	Lys	Ala	Phe	Ser	Gln	Thr	Ser
		275					280					285			
His	Leu	Thr	GIn	His	Gln	Arg	He	His	Ser	Gly	Glu	Thr	Pro	Tyr	Ala
	290					295					300				
Cys	Pro	Val	Cys	G1y	Lys	Ala	Phe	Arg	His	Ser	Ser	Ser	Leu	Val	Arg
305					310					315					320
His	Gln	Arg	He	His	Thr	Ala	Glu	Lys	Ser	Phe	Arg	Cys	Ser	Glu	Cys
				325					330					335	
Gly	Lys	Ala	Phe	Ser	His	$G_{\ell}^{1}y$	Ser	Asn	Leu	Ser	Gln	His	Arg	Lys	lle
			340					345					350		
llis	Ala	Gly	Gly	Arg	Pro	Tyr	Ala	Cys	Ala	Gln	Cys	Gly	Arg	Arg	Phe
		355					360					365			
Cys	Arg	Asn	Ser	His	Leu	Пe	Gln	His	Glu	Arg	Thr	His	Thr	Gly	Glu
	370					375					380				
Lys	Pro	Pho	Val	Cve	A12	Lau	Ara	G1v	Ala	Ala	Dha	Can	Cln	Clv	Sor
	110	1 110	, 611	Cys	M. a	Leu	m g	Ory	ма	MId	THE	261	111.0	Ory	261

Ser Leu Phe Leu His Gln Arg Val His Thr Gly Glu Lys Pro Phe Ala Cys Ala Gln Cys Gly Arg Ser Phe Ser Arg Ser Ser Asn Leu Thr Gln His Gln Leu Leu His Thr Gly Glu Arg Pro Phe Arg Cys Val Asp Cys Gly Lys Gly Phe Ala Lys Gly Ala Val Leu Leu Ser His Arg Arg 11e His Thr Gly Glu Lys Pro Phe Val Cys Thr Gln Cys Gly Arg Ala Phe Arg Glu Arg Pro Ala Leu Leu His His Gln Arg Ile His Thr Thr Glu Lys Thr Asn Ala Ala Ala Pro Asp Cys Thr Pro Gly Pro Gly Phe Leu Gln Gly His His Arg Lys Val Arg Arg Gly Gly Lys Pro Ser Pro Val Leu Lys Pro Ala Lys Val

<210> 4908

<211> 1300

<212> PRT

<213> Homo sapiens

<400> 531

 Met
 Pro
 Pro
 Asp
 Val
 11e
 Arg
 Asp
 Ile
 Leu
 Glu
 Gly
 Val
 Leu
 Arg
 Leu

 Met
 Gly
 Ile
 Phe
 Asp
 Thr
 Ser
 Trp
 Val
 Ser
 Met
 Lys
 Ser
 Phe
 Leu
 Ala

 Lys
 Arg
 Gly
 Val
 Arg
 Glu
 Asp
 Ile
 Ala
 Thr
 Phe
 Asp
 Ala
 Arg
 Asn
 Ile

 Ser
 Lys
 Glu
 Ile
 Asp
 Glu
 Ser
 Val
 Glu
 Glu
 Leu
 Phe
 Lys
 Asn
 Lys

 Ser
 Phe
 Asp
 Pro
 Lys
 Asn
 Ala
 Lys
 Arg
 Ala
 Ser
 Thr
 Ala
 Ala
 Ala

Pro	Leu	Ala	Ala	Trp	Val	Lys	Ala	Asn	Пe	Gln	Tyr	Ser	His	Val	Leu
				85					90					95	
Glu	Arg	He	His	Pro	Leu	Glu	Thr	Glu	Gln	Ala	Gly	Leu	Glu	Ser	Asn
			100					105					110		
Leu	Lys	Lys	Thr	Glu	Asp	Arg	Lys	Arg	Lys	Leu	Glu	Glu	Leu	Leu	Asn
		115					120					125			
Ser	Val	Gly	Gln	Lys	Val	Ser	G]u	Leu	Lys	Glu	Lys	Phe	Gln	Ser	Arg
	130					135					140				
Thr	Ser	Glu	Ala	Ala	Lys	Leu	Glu	Ala	Glu	Val	Ser	Lys	Ala	Gln	Glu
145					150					155					160
Thr	He	Lys	Λla	Ala	Glu	Val	Leu	lle	Asn	Gln	Leu	Asp	Arg	Glu	His
				165					170					175	
Lys	Arg	Trp	Asn	Ala	Gln	Val	Val	Glu	lle	Thr	Glu	Glu	Leu	Ala	Thr
			180					185					190		
Leu	Pro	Lys	Arg	Ala	Gln	Leu	Ala	Ala	Λla	Phe	lle	Thr	Tyr	Leu	Ser
		195					200					205			
Ala	Ala	Pro	Glu	Ser	Leu	Arg	Lys	Thr	Cys	Leu	Glu	Glu	Trp	Thr	Lys
	210					215					220				
Ser	Ala	Gly	Leu	Glu	Lys	Phe	Asp	Leu	Arg	Arg	Phe	Leu	Cys	Thr	Glu
225					230					235					240
Ser	Glu	Gln	Leu	He	Trp	Lys	Ser	Glu	Gly	Leu	Pro	Ser	Asp	Asp	Leu
				245					250					255	
Ser	lle	Glu	Asn	Ala	Leu	Val	He	Leu	Gln	Ser	Arg	Val	Cys	Pro	Phe
			260					265					270		
Leu	lle	Asp	Pro	Ser	Ser	Gln	Ala	Thr	Glu	Trp	Leu	Lys	Thr	His	Leu
		275					280					285			
Lys	Asp	Ser	Arg	Leu	G]u	Val	He	Asn	Gln	Gln	Asp	Ser	Asn	Phe	lle
	290					295					300				
Thr	Ala	Leu	Glu	Leu	Ala	Val	Arg	Phe	Gly	Lys	Thr	Leu	He	He	Gln
305					310					315					320
Glu	Met	Asp	Gly	Val	Glu	Pro	Val	Leu	Tyr	Pro	Leu	Leu	Arg	Arg	Asp
				325					330					335	
Leu	Val	Ala	Gln	Gly	Pro	Arg	Tyr	Val	Val	Gln	He	Gly	Asp	Lys	He
			340					345					350		
lle	Asp	Tyr	Asn	Glu	Glu	Phe	Arg	Leu	Phe	Leu	Ser	Thr	Arg	Asn	Pro
		355					360					365			

Asn	Pro	Phe	He	Pro	Pro	Asp	Ala	Ala	Ser	He	Val	Thr	Glu	Val	Asn
	370					375					380				
Phe	Thr	Thr	Thr	Arg	Ser	Gly	Leu	Arg	Gly	Gln	Leu	Leu	Ala	Leu	Thr
385					390					395					400
Пе	Gln	His	Glu	Lys	Pro	Asp	Leu	Glu	Glu	Gln	Lys	Thr	Lys	Leu	Leu
				405					410					415	
Gln	Gln	Glu	Glu	Asp	Lys	Lys	Пе	Gln	Leu	Ala	Lys	Leu	Glu	Glu	Ser
			420					425					430		
Leu	Leu	Glu	Thr	Leu	Ala	Thr	Ser	Gln	Gly	Asn	He	Leu	Glu	Asn	Lys
		435					440					445			
Asp	Leu	He	Glu	Ser	Leu	Asn	Gln	Thr	Lys	Ala	Ser	Ser	Ala	Leu	11e
	450					455					460				
Gln	Glu	Ser	Leu	Lys	Glu	Ser	Tyr	Lys	Leu	Gln	He	Ser	Leu	Asp	Gln
465					470					475					480
Glu	Arg	Asp	Ala	Tyr	Leu	Pro	Leu	Ala	Glu	Ser	Ala	Ser	Lys	Met	Tyr
				485					490					495	
Phe	lle	lle	Ser	Asp	Leu	Ser	Lys		Asn	Asn	Met	Tyr	Arg	Phe	Ser
			500					505					510		
Leu	Ala		Phe	Leu	Arg	Leu		Gln	Arg	Ala	Leu		Asn	Lys	Gln
		515					520					525	_		
Asp		Glu	Asn	Thr	Glu	Gln	Arg	He	Gln	Ser		He	Ser	Ser	Leu
	530			_		535				_	540				
	His	Met	Val	Tyr		Tyr	He	Cys	Arg		Leu	Phe	Lys	Ala	
545	,	14	DI		550		101	17 7		555			10	61	560
GIn	Leu	Met	Phe		Leu	His	Phe	Val		61 y	Met	HIS	Pro		Leu
DL.	C1	C1	Λ	565	т	Α	ть	DI	570	C1	V - 1	V = 1	V 1	575	A
Pne	GIN	610		610	тр	Asp	inr		Inr	GIŸ	vai	vai		GIŸ	Asp
Mad	1	A	580	۸۱	A ===	C	C1 m	585	Lua	11.	A 20 ~	Aan	590	1	Duo
we t	Leu	595	LyS	ма	nsp	Ser	600	0111	Lys	116	MIG	605		Leu	110
Sor	Trn		Aen	G1n	Glu	Arg		Trn	Ala	Va 1	Λla		Lau	lvc	Ha
361	610	110	изр	OIII	Olu	615	261	пр	MIG	v a 1	620	1111	Leu	rys	110
Ala		Pro	Ser	Lou	Tyr	Gln	Thr	Leu	Cve	Phe		Aen	Ala	Ala	Leu
625	Lou	130	J	cu	630	OIH		12 C CI	0,0	635	O i u	пор	,,,,,,		640
	Arø	Thr	Tvr	Tvr		Asn	Ser	Met	Cvs		Gln	Glu	Phe	Pro	
12	0		- J -	645					650		• •			655	

т1.	1	A 1 -	1	1	V - 1	C	1	DL -	C1	C1	т1.	1	V = 1	V - 1	C1.
116	Leu	Ala		Lys	val	ser	Leu		GIn	GIn	116	Leu		val	GIn
A10	Lou	Ara	660 Pro	Acr	Ara	Lov	Gla	665 Ser	410	Mot	ΛΙο	Lov	670	A10	Cvc
N I M	Leu	675	LTO	кѕр	wrg	Leu	G1n 680	ser.	ита	met	MIA	685	гие	MIB	Cys
Lve	Thr		Clv	Lou	Lvc	Clu	Val	Sor	Pro	Lou	Pro		Acn	Lou	lve
Lys	690	Leu	Gry	ren	Lys	695	vai	361	110	Leu	700	Leu	изп	Leu	Lys
Ara		Tyr	lve	Glu	The		Glu	He	Glu	Pro		Leu	He	He	He
705	LÇU	1) 1	123.3	Olu	710	Lea	Q1 ti	.1.1.0	010	715	.110	Lea	.110	110	720
	Pro	G1 v	Ala	Asn		Ser	Gln	Glu	Len		Glu	Leu	Ala	Asn	
001	110	Oly	7114	725	110	501	0111	oru	730	0111	014	Вод		735	,,,,
Glu	Arg	Ser	Glv		Cvs	Tvr	His	Gln		Ala	Met	Glv	Gln		Gln
	0		740			•		745				,	750	,	
Ala	Asp	Leu	Ala	He	G]n	Met	Leu	Lys	Glu	Cys	Ala	Arg	Asn	G1 y	Λsp
		755					760					765			
Trp	Leu	Cys	Leu	Lys	Asn	Leu	His	Leu	Val	Val	Ser	Trp	Leu	Pro	Val
	770					775					780				
Leu	Glu	Lys	Glu	Leu	Asn	Thr	Leu	Gln	Pro	Lys	Asp	Thr	Phe	Arg	Leu
785					790					795					800
Trp	Leu	Thr	Ala	Glu	Val	His	Pro	Asn	Phe	Thr	Pro	lle	Leu	Leu	Gln
				805					810					815	
Ser	Ser	Leu	Lys	He	Thr	Tyr	Glu	Ser	Pro	Pro	Gly	Leu	Lys	Lys	Asn
			820					825					830		
Leu	Met	Arg	Thr	Tyr	Glu	Ser	Trp	Thr	Pro	Glu	Gln	He	Ser	Lys	Lys
		835					840					845			
Asp		Thr	llis	Arg	Ala		Ala	Leu	Phe	Ser		Ala	Trp	Phe	His
	850					855				_	860				
	Ala	Cys	Gln	Glu		Arg	Asn	Tyr	He		G1n	G1 y	Trp	Thr	
865		0.1	51		870			,		875	0.1	m			880
Phe	Tyr	Glu	Phe		Leu	Ser	Asp	Leu		Ala	GIy	Tyr	Asn		He
			DI	885	61	. 1			890	C1	т	C.1	DI	895	
Asp	Arg	Leu		Asp	GIY	Ala	Lys		Val	GIN	Lrp	Glu		vai	HIS
C1.	1	1	900	Λ	Д1	11.	т	905	C1.	Λ	11.	Λ	910	т	D1
оту	ren		oru	ASN	MIA	116	Tyr	GIŸ	ыу	лгg	911,	925	ASII	ryr	rne
٨٥٠	Lau	915	Vol	Lau	Cl-	San	920	Lau	1,,,,	C1.	Dlsa		A	Son	S ~ **

	930					935					940				
Val	He	Asp	Val	Phe	Asn	Gln	Arg	۸sn	Lys	Lys	Ser	He	Phe	Pro	Tyr
945					950					955					960
Ser	Val	Ser	Leu	Pro	Gln	Ser	Cys	Ser	lle	Leu	Asp	Tyr	Arg	Ala	Val
				965					970					975	
He	Glu	Lys	lle	Pro	Glu	Asp	Asp	Lys	Pro	Ser	Phe	Phe	Gly	Leu	Pro
			980					985					990		
Ala	Asn	Ile	Ala	Arg	Ser	Ser	Gln	Arg	Met	lle	Ser	Ser	Gln	Val	He
		995					1000					1005			
Ser	Gln	Leu	Arg	Ile	Leu	Gly	Arg	Ser	He	Thr	Ala	Gly	Ser	Lys	Phe
]	010]	1015					1020				
Asp	Arg	Glu	He	Trp	Ser	Asn	Glu	Leu	Ser	Pro	Val	Leu	Asn	Leu	Trp
1025	5				1030					1035					1040
Lys	Lys	Leu	Asn	Gln	Asn	Ser	Asn	Leu	He	His	G1n	Lys	Val	Pro	Pro
]	045					1050					1055	
Pro	Asn	Asp	Arg	Gln	Gly	Ser	Pro	lle	Leu	Ser	Phe	lle	lle	Leu	Glu
			1060					1065					1070		
Gln	Phe	Asn	Ala	lle	Arg	Leu	Val	Gln	Ser	Val	His	Gln	Ser	Leu	Ala
		1075					1080					1085			
Ala	Leu	Ser	Lys	Val	lle	Arg	Gly	Thr	Thr	Leu	Leu	Ser	Ser	Glu	Va]
]	1090					1095					1100				
Gln	Lys	Leu	Ala	Ser	Ala	Leu	Leu	Asn	Gln	Lys	Cys	Pro	Leu	Ala	Trp
1109	5				1110					1115					1120
Gln	Ser	Lys	Trp	Glu	Gly	Pro	Glu	Asp	Pro	l.eu	Gln	Tyr	Leu	Arg	Gly
				1125					1130					1135	
Leu	Val	Ala	Arg	Ala	Leu	Ala	lle	Gln	Asn	Trp	Val	Asp	Lys	Ala	Glu
			1140					1145					1150		
Lys			Leu	Leu	Ser			Leu	Asp	Leu		Glu	Leu	Phe	His
		1155					1160					1165			
		Thr	Phe	Leu			Leu	Arg	GIn			Ala	Arg	Ala	Val
	1170	_				1175	_				1180	_	-		
		Ser	Val			Leu	Lys	Phe			Ser	Trp	Lys		
1185					1190					1195					1200
Leu	GIn	Glu			Leu	GIn	He			Ser	Gly	Leu			G.Lu
				1205					1210					1215	
61	Cvc	C	101 -	Λ	C^{1}	Λ .	C^{1}	1	C	C1.	Λ	C1	1	Λ	C

1225 1230 1220 Pro Ser Val Ser Ser Val Leu Pro Cys Phe Met Gly Trp 11e Pro Gln 1240 Asp Ala Cys Gly Pro Tyr Ser Pro Asp Glu Cys Ile Ser Leu Pro Val 1250 1255 1260 Tyr Thr Ser Ala Glu Arg Asp Arg Val Val Thr Asn Ile Asp Val Pro 1270 1275 Cys Gly Gly Asn Gln Asp Gln Trp Ile Gln Cys Gly Ala Ala Leu Phe 1290 Leu Lys Asn Gln 1300 <210> 4909 <211> 944 <212> PRT <213> Homo sapiens <400> 532 Met Gly Ser Asp Gly 11e Leu Arg Leu Ser Thr Ser Ala Leu Asn Asn 1 5 10 15 Glu Phe Phe Ala Tyr Ala Ala Gln Gly Trp Lys Gln Arg Leu Ala Glu 25 Gly Glu Phe Thr Pro Glu Met Gln Leu Arg Ile Arg Gln Glu Ile Glu 35 40 45 Lys Glu Lys Lys Thr Glu Pro Trp Lys Glu Lys Phe Phe Glu Arg Phe 55 Tyr Gly Glu Lys Leu Gly Met Ser Arg Glu Glu Ser Val Lys Leu Thr 70 75 Thr Gly Pro Asn Asn Ala Gly Ala Gln Ser Ser Ser Ser Cys Gly Thr 85 90 95 Ser Gly Leu Pro Val Ser Ala Gln Thr Ala Leu Ala Glu Gln Gln Pro 105 Lys Ser Met Lys Ser Pro Ala Ser Pro Glu Pro Gly Phe Cys Ala Thr 115 . 120

Leu Cys Pro Met Val Glu lle Pro Pro Lys Asp lle Met Ala Glu Leu

	130					135					140				
Glu	Ser	Glu	Asp	He	Leu	lle	Pro	Glu	Glu	Ser	Val	Ile	Gln	Glu	Glu
145					150					155					160
He	Ala	Glu	Glu	Val	Glu	Thr	Ser	Пе	Cys	Glu	Cys	Gln	Asp	Glu	Asn
				165					170					175	
His	Lys	Thr	He	Pro	Glu	Phe	Ser	Glu	Glu	Ala	Glu	Ser	Leu	Thr	Asn
			180					185					190		
Ser	His	Glu	Glu	Pro	Gln	lle	Ala	Pro	Pro	Glu	Asp	Asn	Leu	Glu	Ser
		195					200					205			
Cys	Val	Met	Met	Asn	Asp	Val	Leu	Glu	Thr	Leu	Pro	His	lle	Glu	Val
	210					215					220				
Lys	lle	Glu	Gly	Lys	Ser	Glu	Ser	Pro	Gln	Glu	Glu	Met	Thr	Val	Val
225					230					235					240
lle	Asp	Gln	Leu	Glu	Val	Cys	Asp	Ser	Leu	He	Pro	Ser	Thr	Ser	Ser
				245					250					255	
Met	Thr	His	Val	Ser	Asp	Thr	Glu	His	Lys	Glu	Ser	Glu	Thr	Ala	Val
			260					265					270		
Glu	Thr	Ser	Thr	Pro	Lys	Ile	Lys	Thr	Gly	Ser	Ser	Ser	Leu	Glu	Gly
		275					280					285			
Gln	Phe	Pro	Asn	Glu	Gly	Ile	Ala	Ile	Asp	Met	Glu	Leu	Gln	Ser	Asp
	290					295					300				
Pro	Glu	Glu	Gln	Leu	Ser	Glu	Asn	Ala	Cys	11e	Ser	Glu	Thr	Ser	Phe
305					310					315					320
Ser	Ser	Glu	Ser	Pro	Glu	G1 y	Ala	Cys	Thr	Ser	Leu	Pro	Ser	Pro	G1 y
				325					330					335	
Gly	Glu	Thr	Gln	Ser	Thr	Ser	Glu	Glu	Ser	Cys	Thr	Pro	Ala	Ser	Leu
			340					345					350		
Glu	Thr	Thr	Phe	Cys	Ser	Glu	Val	Ser	Ser	Thr	Glu	Asn	Thr	Asp	Lys
		355					360					365			
Tyr	Asn	Gln	Arg	Asn	Ser	Thr	Asp	Glu	Asn	Phe	His	Ala	Ser	Leu	Met
	370					375					380				
Ser	Glu	He	Ser	Pro	lle	Ser	Thr	Ser	Pro	Glu	He	Ser	Glu	Ala	Ser
385					390					395					400
Leu	Met	Ser	Asn	Leu	Pro	Leu	Thr	Ser	Glu	Ala	Ser	Pro	Val	Ser	Asn
				405					410					415	
Leu	Pro	Leu	Thr	Ser	G] u	Thr	Ser	Pro	Met	Ser	Asp	Leu	Pro	Leu	Thr

			420					425					430		
Ser	Lys	Thr	Ser	Ser	Val	Ser	Ser	Met	Leu	Leu	Thr	Ser	Glu	Thr	Thr
		435					440					445			
Phe	Val	Ser	Ser	Leu	Pro	Leu	Pro	Ser	Glu	Thr	Ser	Pro	He	Ser	Asn
	450					455					460				
Ser	Ser	Пе	Λsn	Glu	Arg	Met	Ala	His	Gln	Gln	Arg	Lys	Ser	Pro	Ser
465					470					475					480
Val	Ser	Glu	Glu	Pro	Leu	Ser	Pro	Gln	Lys	Asp	Glu	Ser	Ser	Ala	Thr
				485					490					495	
Ala	Lys	Pro	Leu	Gly	Glu	Asn	Leu	Thr	Ser	Gln	Gln	Lys	Asn	Leu	Ser
			500					505					510		
Asn	Thr	Pro	Glu	Pro	lle	Ile	Met	Ser	Ser	Ser	Ser	lle	Ala	Pro	Glu
		515					520					525			
Ala	Phe	Pro	Ser	Glu	Asp	Leu	His	Asn	Lys	Thr	Leu	Ser	Gln	Gln	Thr
	530					535					540				
Cys	Lys	Ser	His	Val	Asp	Thr	Glu	Lys	Pro	Tyr	Pro	Ala	Ser	He	Pro
545					550					555					560
Glu	Leu	Ala	Ser	Thr	Glu	Met	Ile	Lys	Val	Lys	Asn	His	Ser	Val	Leu
				565					570					575	
Gln	Arg	Thr	Glu	Lys	Lys	Val	Leu	Pro	Ser	Pro	Leu	Glu	Leu	Ser	Val
			580					585					590		
Phe	Ser	Glu	Gly	Thr	Asp	Asn	Lys	Gly	Asn	Glu	Leu	Pro	Ser	Ala	Lys
		595					600					605			
Leu	Gln	Asp	Lys	Gln	Tyr	lle	Ser	Ser	Val	Asp	Lys	Ala	Pro	Phe	Ser
	610					615					620				
Glu	Gly	Ser	Arg	Asn	Lys	Thr	His	Lys	Gln	Gly	Ser	Thr	Gln	Ser	Arg
625					630					635					640
Leu	Glu	Thr	Ser	His	Thr	Ser	Lys	Ser	Ser	Glu	Pro	Ser	Lys	Ser	Pro
				645					650					655	
Asp	Gly	He	Arg	Asn	Glu	Ser	Arg	Asp	Ser	Glu	He	Ser	Lys	Arg	Lys
			660					665					670		
Thr	Ala	Glu	Gln	His	Ser	Phe	Gly	He	Cys	Lys	Glu	Lys	Arg	Ala	Arg
		675					680					685			
lle		Asp	Asp	G1n	Ser		Arg	Asn	lle	Ser	Ser	Ser	Ser	Pro	Pro
	690					695					700				
G1n	Lvc	Glu	Gla	Pro	Pro	Ara	Glin	GL_{1}	Pro	Ara	$V_2 1$	Pro	Pro	Lau	lve

705					710					715					720
Пe	Gln	Leu	Ser	Lys	He	Gly	Pro	Pro	Phe	lle	lle	Lys	Ser	Gln	Pro
				725					730					735	
Va]	Ser	Lys	Pro	Glu	Ser	Arg	Ala	Ser	Thr	Ser	Thr	Ser	Val	Ser	Gly
			740					745					750		
Gly	Arg	Asn	Thr	Gly	Ala	Arg	Thr	Leu	Ala	Asp	He	Lys	Ala	Arg	Ala
		755					760					765			
Gln	Gln	Ala	Arg	Ala	Gln	Arg	Glu	Ala	Ala	Ala	Ala	Ala	Ala	Val	Ala
	770					775					780				
Ala	Ala	Ala	Ser	Ile	Val	Ser	Gly	Ala	Met	Gly	Ser	Pro	Gly	Glu	Gly
785					790					795					800
Gly	Lys	Thr	Arg	Thr	Leu	Ala	His	lle	Lys	Glu	Gln	Thr	Lys	Ala	Lys
				805					810					815	
Leu	Phe	Ala	Lys	His	Gln	Ala	Arg	Ala	His	Leu	Phe	Gln	Thr	Ser	Lys
			820					825					830		
Glu	Thr	Arg	Leu	Pro	Pro	Pro	Leu	Ser	Ser	Lys	Glu	Gly	Pro	Pro	Asn
		835					840					845			
Leu	Glu	Val	Ser	Ser	Thr	Pro	Glu	Thr	Lys	Met	Glu	Gly	Ser	Thr	G1 y
	850					855					860				
Val	He	He	Val	Asn	Pro	Asn	Cys	Arg	Ser	Pro	Ser	Asn	Lys	Ser	Ala
865					870					875					880
His	Leu	Arg	Glu	Thr	Thr	Thr	Val	Leu	Gln	Gln	Ser	Leu	Asn	Pro	Ser
				885					890					895	
Lys	Leu	Pro	Glu	Thr	Ala	Thr	Asp	Leu	Ser	Val	His	Ser	Ser	Asp	Glu
			900					905					910		
Asn	Ile	Pro	Val	Ser	His	Leu		Glu	Lys	lle	Val		Ser	Thr	Ser
		915					920					925			
Ser		Asn	Ser	Ser	Val	Pro	Met	Leu	Phe	Asn	Lys	Asn	Ser	Val	Pro
	930					935					940				

<210> 4910

<211> 1161

<212> PRT

<213> Homo sapiens

<400)> 53	33													
Met	Cys	Ser	Thr	Pro	Gly	Met	Pro	Ala	Pro	Gly	Ala	Ser	Leu	Ala	Leu
1				5					10					15	
Arg	Val	Ser	Phe	Val	Asp	Val	His	Pro	Asp	Val	He	Pro	Val	G1n	Leu
			20					25					30		
Trp	Gly	Leu	Val	Gly	Glu	Arg	Arg	Gly	Glu	Tyr	Leu	Arg	Leu	Ser	Arg
		35					40					45			
Glu	He	Gln	Glu	Ala	Ala	Ala	Thr	Arg	Gly	Gln	Trp	Ala	Leu	Gly	Ser
	50					55					60				
Ala	Ser	Ala	Ser	Pro	Gly	Glu	Leu	Cys	Leu	Val	Gln	Val	Gly	Leu	Leu
65					70					75					80
Trp	His	Arg	Cys	Arg	Val	Val	Ser	Arg	Gln	Ala	Gln	Glu	Ser	Arg	Val
				85					90					95	
Phe	Leu	Leu	Asp	Glu	Gly	Arg	Thr	He	Thr	Ala	Gly	Ala	Gly	Ser	Leu
			100					105					110		
Ala	Pro	Gly	Arg	Arg	Glu	Phe	Phe	Asn	Leu	Pro	Ser		Val	Leu	Gly
		115					120					125			
Cys		Leu	Ala	Gly	Leu		Pro	Ala	Gly	Cys		Ala	Gly	Ser	Gly
	130					135					140				
	Pro	Pro	Gln	His		Pro	Ala	Asp	Ala		Asp	Phe	Leu	Ser	
145			_		150			_		155					160
Leu	GIn	Gly	Lys		Val	His	Gly	Cys		Leu	Asp	Val	Leu		Leu
			., .	165		0.1	., .	n	170		101	0.1	0.1	175	
HIS	Arg	Leu		Leu	Leu	Glu	Val		Asp	Val	Phe	GIn		Met	Arg
Cl	1	C1 -	180	A 7	Α.	Δ.	V. 1	185 D	Α	C	1	DI	190	C	1
61 u	Leu		ı.eu	Ala	Arg	Arg	Val		ASP	ser	Leu	205	Arg	Ser	Leu
ر ده ا	C1	195	Turn	1	Than	41.	200		110	Son	Vol		Can	C1	Vo 1
Leu		MIG	1 7 1	Leu	m		Ala	1111	мта	261	220	GIY	sei	GIY	val
Dro	210	Lou	Sor	Ara	Vo.1	215 Pro	Leu	1 00	G1 ₁₀	Lvc		Pro	C1 _v	Lou	Acn
225	vai	Leu	361	ni g	230	110	Leu	Lys	0111	235	OIII	110	Oly	Leu	240
	Pho	Tyr	Pro	Gln		Gln	Leu	Glv	Val		Glu	Λla	Val	Val	
1 y 1	1 110	1)1	. 10	245	Leu	0111	i.cu	013	250	1111	Olu	111 a	, a i	255	110
Thr	Gln	Val	Cve		Pro	Hic	Arg	Tle		Cvs	Gln	PII	Arø		Val
1111	0111		260	1113	110	1,13	111 g	265	1113	0,3	OIN	1,00	270	501	
Sor	Gln	Glu		Hic	Δεσ	ا ما	Sor		Ser	Met	Λla	Gln		Tyr	Aro

		275					280					285			
Gly	Ser	Thr	Gly	Thr	Gly	Asp	Glu	۸sn	Ser	Thr	Ser	Ala	Thr	Trp	Glu
	290					295					300				
Glu	Arg	Glu	G1u	Ser	Pro	Asp	Lys	Pro	Gly	Ser	Pro	Cys	Ala	Ser	Cys
305					310					315					320
G1 y	Leu	Asp	G1y	His	Trp	Tyr	Arg	Ala	Leu	Leu	Leu	Glu	Thr	Phe	Arg
				325					330					335	
Pro	Gln	Arg	Cys	Ala	Gln	Val	Leu	His	Val	Asp	Tyr	Gly	Arg	Lys	Glu
			340					345					350		
Leu	Val	Ser	Cys	Ser	Ser	Leu	Arg	Tyr	Leu	Leu	Pro	Glu	Tyr	Phe	Arg
		355					360					365			
Met	Pro	Val	Val	Thr	Tyr	Pro	Cys	Ala	Leu	Tyr	Gly	Leu	Trp	Asp	Gly
	370					375					380				
Gly	Arg	Gly	Trp	Ser	Arg	Ser	Gln	Val	Gly	Asp	Leu	Lys	Thr	Leu	lle
385					390					395					400
Leu	Gly	Lys	Ala	Val	Asn	Ala	Lys	Πe	Glu	Phe	Tyr	Cys	Ser	Phe	Glu
				405					410					415	
His	Val	Tyr	Tyr	Val	Ser	Leu	Tyr	Gly	Glu	Asp	Gly	lle	Asn	Leu	Asn
			420					425					430		
Arg	Val	Phe	Gly	Val	Gln	Ser	Cys	Cys	Leu	Ala	Asp	Arg	Val	Leu	Gln
		435					440					445			
Ser	Gln	Ala	T <u></u> hr	Glu	Glu	Glu	Glu	Pro	Glu	Thr	Ser	Gln	Ser	Gln	Ser
	450					455					460				
Pro	Ala	Glu	Glu	Val	Asp	Glu	Glu	lle	Ser	Leu	Pro	Ala	Leu	Arg	Ser
465					470					475					480
Ile	Arg	Leu	Lys	Met	Asn	Ala	Phe	Tyr	Asp	Ala	Gln	Val	Glu	Phe	Val
				485					490			•		495	
Lys	Asn	Pro	Ser	Glu	Phe	Trp	He	Arg	Leu	Arg	Lys	His	Asn	Val	Thr
			500					505					510		
Phe	Ser	Lys	Leu	Met	Arg	Arg	Met	Cys	G]y	Phe	Tyr	Ser	Ser	Ala	Ser
		515					520					525			
Lys	Leu	Asp	Gly	Val	Val	Leu	Lys	Pro	Glu	Pro	Asp	Asp	Leu	Cys	Cys
	530					535					540				
Val	Lys	Trp	Lys	Glu	Asn	Gly	Tyr	Tyr	Arg	Ala	Пe	Val	Thr	Lys	Leu
545					550					555					560
Asp	Asp	Lys	Ser	Val	Asp	Val	Phe	Leu	Val	Asp	Arg	Gly	Asn	Ser	Glu

					565					570					575	
As	sn	Val	Asp	Trp	Tyr	Asp	Val	Arg	Met	Leu	Leu	Pro	Gln	Phe	Arg	Gln
				580					585					590		
Le	eu	Pro	Пe	Leu	Ala	Val	Lys	Cys	Thr	Leu	Ala	Asp	Пę	Trp	Pro	Leu
			595					600					605			
G1	lу	Lys	Thr	Trp	Ser	Gln	Glu	Ala	Val	Ser	Phe	Phe	Lys	Lys	Thr	Val
		610					615					620				
Le	eu	His	Lys	Glu	Leu	Val	Ile	His	He	Leu	Asp	Lys	Gln	Asp	His	Gln
62	25					630					635					640
Ty	/r	Val	Ile	Glu	He	Leu	Asp	Glu	Ser	Arg	Thr	Gly	Glu	Glu	Asn	Ile
					645					650					655	
Se	er	Lys	Val	He	Ala	Gln	Ala	Gly	Tyr	Ala	Lys	Tyr	Gln	Glu	Phe	G] u
				660					665					670		
Tł	ır	Lys	Glu	Asn	He	Leu	Val	Asn	Ala	His	Ser	Pro	Gly	His	Val	Ser
			675					680					685			
٨s	sn	His	Phe	Thr	Thr	Glu	Ser	Asn	Lys	He	Pro	Phe	Ala	Lys	Thr	Gly
		690					695					700				
G 3	lu	Gly	Glu	Gln	Lys	Ala	Lys	Arg	Glu	Asn	Lys	Thr	Thr	Ser	Val	Ser
)5					710					715					720
Ly	/S	Ala	Leu	Ser	Asp	Thr	Thr	Val	Val	Thr	Asn	Gly	Ser	Thr	Glu	Leu
					725					730					735	
Va	a l	Val	Gln		Lys	Val	Lys	Arg		Ser	Val	Tyr	Phe		Leu	Met
				740					745					750		
G]	ln	Asn		Leu	Glu	He	Lys		Gly	Ser	Ser	Ser		Gly	Glu	Leu
			755	_				760				_	765	~ "		_
G.	lu		Gly	Ser	Thr	Val			Arg	Val	Ser			Glu	Asn	Pro
0.1		770	13.1		0	a 1	775					780				m i
		lyr	Phe	Irp	Cys	Gln	Leu	lhr	Arg	Asn		GIn	GIÿ	Leu	Lys	
	35				7.1	790	T	Tr.	0	,	795	3 11	. 1	• 1	D	800
Le	eu	Met	Ser	Asp		Gln	lyr	iyr	Cys		Asn	lhr	Ala	Ala		HIS
C1	١	Δ	Α	Tl	805	A 1 -	C	1	A 1 =	810	Α	Tl	V. 1	A	815	C1
G.	ın	Arg	Asn		Leu	Ala	Cys	Leu		Lys	Arg	Inr	vai		Arg	GIN
т		Carr	A	820	1	71.	C	C1-	825	C1	C	V. I	C1.	830	V = 1	Λ =:
1.1	гþ	ser		Ala	Leu	Ile	ser		116	oin	ser	val		n1s	val	ASN
			835					840					845			
v	. 1	Tl	D1	Val	Λ	Т	C1	Λ	Λ	C^{1}	14 - 4	Vi - 1	C	V ~ 1	44.0	Λ

	850					855					860				
Πle	Tyr	Ser	He	Ser	Glu	Glu	Phe	Leu	Lys	Val	Lys	Ala	Gln	Ala	Phe
865					870					875					880
Arg	Cys	Ser	Leu	Tyr	Asn	Leu	lle	Gln	Pro	Val	Gly	Gln	Asn	Pro	Phe
				885					890					895	
Va]	Trp	Asp	Val	Lys	Ala	Пе	Gln	Ala	Phe	Asn	Glu	Phe	lle	Asp	Asn
			900					905					910		
Ala	Trp	Gln	Lys	Asn	Leu	Glu	Leu	Lys	Cys	Thr	He	Phe	Ala	Leu	Ala
		915					920					925			
Ser	lle	Asn	Glu	Glu	Leu	Р́hе	Asn	He	Val	Asp	Leu	Leu	Thr	Pro	Phe
	930					935					940				
Gln	Ser	Ala	Cys	His	Phe	Leu	Val	G]u	Lys	Arg	Leu	Ala	Arg	Pro	Va]
945					950					955					960
Lys	Leu	Gln	Lys	Pro	Leu	Glu	Ser	Ser	Val	Gln	Leu	His	Ser	Tyr	Phe
				965					970					975	
Tyr	Ser	Thr	His	Asp	Met	Lys	He	Gly	Ser	Glu	Glu	Leu	Val	Tyr	lle
			980					985					990		
Thr	His	lle	Asp	Asp	Pro	Trp	Thr	Phe	Tyr	Cys	Gln	Leu	Ala	Arg	Asn
		995]	1000					1005			
Ala	Asn	He	Leu	Glu	Gln	Leu	Ser	Cys	Ser			Gln	Leu	Ser	Lys
	1010					1015					1020				
Val	Leu	Leu	Asn	Leu	Lys	Thr	Ser	Pro	Leu	Asn	Pro	Gly	Thr	Leu	Cys
1025					1030					1035					1040
Leu	Ala	Lys			Asp	G1 y	Asn			Arg	Gly	lle			Glu
				1045					1050					1055	
Lys	Glu			Lys	Val	Phe			Asp	Phe	Gly			Tyr	Val
			1060					1065					1070		
Val			Asp	Asp	Leu	Leu		He	Pro	Ser			lyr	Asp	Val
,		1075	D.	14 .	61		1080		C	C		085		11	D
		Leu	Pro	Met		Ala	vai	Arg	Cys			Ser	Asp	116	Pro
	1090	11.	D	C1		1095	V - 1	V - 1	Т		1100	C1	Tl	31.	1
		He	Pro			Val	val	val			GIn	Glu	Inr		
110		Son	Lau		1110	1	Ve1	Ve 1		1115	۸۵.	Dance	A 655		1120 The
asp	Lys	ser	Leu	LyS	n18	Leu	val	1 15 4	n i a	LyS	nsp	LLO	nsp	olb	inr
				1195					1130					1135	
Leu	He	116		1125	Tyr	Glv	Aen		1130	Gla	Πa	Ser	_	1135 Ser	116

Asn Lys Lys Leu Gly Leu Leu Ser Tyr <210> 4911 <211> 762 <212> PRT <213> Homo sapiens <400> 534 Met Gly Asp Leu Lys Ser Gly Phe Glu Glu Val Asp Gly Val Arg Leu Gly Tyr Leu Ile Ile Lys Gly Lys Gln Met Phe Ala Leu Ser Gln Val Phe Thr Asp Leu Lys Asn Ile Pro Arg Thr Thr Val His Lys Arg Met Asp His Leu Lys Val Lys Lys His His Cys Asp Leu Glu Glu Leu Arg Lys Leu Lys Ala Ile Asn Ser Ile Ala Phe His Ala Ala Lys Cys Thr Leu lle Ser Arg Glu Asp Val Glu Ala Leu Tyr Thr Ser Cys Lys Thr Glu Arg Val Leu Lys Thr Lys Arg Arg Arg Val Gly Arg Ala Leu Ala Thr Lys Ala Pro Pro Pro Glu Arg Ala Ala Ala Ser Pro Arg Pro Gly Phe Trp Lys Asp Lys His Gln Leu Trp Arg Gly Leu Ser Gly Ala Ala Arg Pro Leu Pro 11e Ser Ala Gln Ser Gln Arg Pro Gly Ala Ala Ala Ala Arg Pro Ala Ala Ilis Leu Pro Gln Ile Phe Ser Lys Tyr

Pro Gly Ser His Tyr Pro Glu lle Val Arg Ser Pro Cys Lys Pro Pro

Leu	Asn	Tyr	Glu	Thr	Ala	Pro	Leu	Gln	Gly	Asn	Tyr	Val	Ala	Phe	Pro
		195					200					205			
Ser	Asp	Pro	Ala	Tyr	Phe	Arg	Ser	Leu	Leu	Cys	Ser	Lys	His	Pro	Λla
	210					215					220				
Ala	Λla	Ala	Ala	Gly	Лlа	Thr	Cys	Leu	Glu	Arg	Phe	His	Leu	Val	Λsn
225					230					235					240
Gly	Phe	Cys	Pro	Pro	Pro	His									
				245					250					255	
His	His	His	His	His	Arg	Ala	Gln	Pro	Pro	Gln	Gln	Ser	His	His	Pro
			260					265					270		
Pro	His	His	His	Arg	Pro	Gln	Pro	His	Leu	Gly	Ser	Phe	Pro	Glu	Ser
		275					280					285			
Cys	Ser	Ser	Asp	Ser	Glu	Ser	Ser	Ser	Tyr	Ser	Asp	His	Ala	Ala	Asn
	290					295					300				
Asp	Ser	Asp	Phe	Gly	Ser	Ser	Leu	Ser	Ser	Ser	Ser	Asn	Ser	Val	Ser
305					310					315					320
Ser	Glu	Glu	Glu	Glu	Glu	Glu	Gly	Glu							
				325					330					335	
Glu	Glu	Glu	Glu	Gly	Gly	Ser	Gly	Ala	Ser	Asp	Ser	Ser	Glu	Val	Ser
			340					345					350		
Ser	Glu	Glu	Glu	Asp	Ser	Ser	Thr	Glu	Ser	Asp	Ser	Ser	Ser	Gly	Ser
		355					360					365			
Ser	Gln	Val	Ser	Val	Gln	Ser	He	Arg	Phe	Arg	Arg	Thr	Ser	Phe	Cys
	370					375					380				
Lys	Pro	Pro	Ser	Val	Gln	Ala	Gln	Ala	Asn	Phe	Leu	Tyr	His	Leu	Ala
385					390					395					400
Ser	Ala	Ala	Ala	Ala	Thr	Lys	Pro	Ala	Ala	Phe	Glu	Asp	Ala	Gly	Arg
				405					410					415	
Leu	Pro	Asp	Leu	Lys	Ser	Ser	Val	Lys	Ala	Glu	Ser	Pro	Ala	Glu	Trp
			420					425					430		
Asn	Leu	Gln	Ser	Trp	Ala	Pro	Lys	Ala	Ser	Pro	Val	Tyr	Cys	Pro	Ala
		435					440					445			
Ser	Leu	Gly	Ser	Cys	Phe	Ala	Glu	He	Arg	Asn	Asp	Arg	Val	Ser	Glu
	450					455					460				
He	Thr	Phe	Pro	His	Ser	Glu	He	Ser	Asn	Ala	Val	Lys	Arg	Thr	Asp
465					470					475					480

Leu	Thr	He	Asn	Cys	Leu	Ala	Glu	Gl y	Ala	Ser	Ser	Pro	Ser	Pro	Lys
				485					490					495	
Thr	Asn	Asn	Аlа 500	Phe	Pro	Gln	Gln	Arg 505	lle	Leu	Arg	Glu	Ala 510	Arg	Lys
Cys	Leu	Gln 515	Thir	Thr	Pro	Thr	Thr 520	His	Cys	Ala	Asp	Asn 525	Asn	Thr	Ile
Ala	Ala 530	Arg	Phe	Leu	Asn	Asn 535	Asp	Ser	Ser	Gly	Ala 540	Glu	Ala	Asn	Ser
G1u 545	Lys	Tyr	Ser	Lys	11e 550	Leu	His	Cys	Pro	Glu 555	Phe	Ala	Thr	Asp	Leu 560
Pro	Ser	Ser	Gln	Thr 565	Asp	Pro	Glu	Val	Asn 570	Ala	Ala	Gly	Ala	Ala 575	Ala
Thr	Lys	Ala	Glu 580	Asn	Pro	Cys	Thr	Asp 585	Thr	Gly	Asp	Lys	Thr 590	Leu	Pro
Phe	Leu	His 595	Asn	11e	Lys	lle	Lys 600	Val	Glu	Asp	Ser	Ser 605	Ala	Asn	Glu
Glu	Tyr 610	Glu	Pro	His	Leu	Phe 615	Thr	Asn	Lys	Leu	Lys 620	Cys	Glu	Cys	Asn
Asp 625	Thr	Lys	Gly	Glu	Phe 630	Tyr	Ser	Val	Thr	Glu 635	Ser	Lys	Glu	Glu	Asp 640
Ala	Leu	Leu	Thr	Thr 645	Ala	Lys	Glu	Gly	Phe 650	Ala	Cys	Pro	Glu	Lys 655	Glu
Thr	Pro	Ser	Leu 660	Asn	Pro	Leu	Ala	G1n 665	Ser	Gln	Gly	Leu	Ser 670	Cys	Thr
Leu	Gly	Ser 675	Pro	Lys	Pro	Glu	Аsр 680	Gly	Glu	Tyr	Lys	Phe 685	Gly	Ala	Arg
Val	Arg 690	Lys	Asn	Tyr	Arg	Thr 695	Leu	Val	Leu	Gly	Lys 700	Arg	Pro	Val	Leu
G1n 705	Thr	Pro	Pro	Val	Lys 710	Pro	Asn	Leu	Lys	Ser 715	Ala	Arg	Ser	Pro	Arg 720
Pro	Thr	Gly	Lys	Thr 725	Glu	Thr	Asn	Glu	Gly 730	Thr	Leu	Asp	Asp	Phe 735	Thr
Val	lle	Asn	Arg 740	Arg	Lys	Lys	Val	Ala 745	Ser	Asn	Val	Ala	Ser 750	Ala	Va]
Lys	Arg	Pro 755	Phe	His	Phe	Met	Ala 760	Asn	Lys						

```
<210> 4912
<211> 619
<212> PRT
<213> Homo sapiens
<400> 535
Met Glu Gly Asp Ala Glu Thr Asp Val Leu Glu Cys Ala Asn Gln Arg
                                     10
                                                          15
Leu Val Ile Ser Glu Thr Asp Gly Glu Ile Leu Thr Pro Gly Trp Asp
                                 25
Thr Gln Asp Arg Met Gly Val Glu Ser Arg Thr Asn lle Gln Glu Leu
                             40
                                                 45
Gly Asn Arg Asn Gln Arg Glu Ala Gly Gly Glu Asn Leu Pro Glu Thr
                         55
Gln Ala His Met Gly Glu Asn Gln Glu Gln Leu Arg Cys Lys lle Asp
                     70
                                         75
Ala Glu Thr Gln Thr Pro Glu Trp Glu Asn Gln Asp Lys Asn Gly Ser
                 85
                                     90
Glu Asp Ala Val Glu Thr Gln Thr Phe Glu Lys Lys Asp Lys Lys Glu
            100
                                105
                                                     110
Ala Gly Glu Glu Asp Gly Glu Glu Ile Gln Ala Gln Gly Leu Gly Lys
                            120
Gln Gly Gln Thr Gly Asp Glu Asn Gly Glu Glu Thr Gln Thr Pro Gln
    130
                        135
Trp Glu Lys Gln Asp Gln Met Lys Gly Asp Ala Asp Val Glu lle Gln
                    150
                                        155
Met Glu Glu Gly Arg Asn Lys Asp Gln Val Gly Gly Gln Asp Ala Ala
                165
                                     170
                                                         175
Gln Thr Gln Ser Cys Gly Arg Glu Asn Val Gly Glu Val Lys Lys Glu
                                185
Asn Ser Val Glu Thr Gln Ala Leu Asp Trp Gly Lys Gln Glu Cys Val
        195
                            200
                                                 205
Gly Asn Gly Asn Val Thr Glu 11e Gln Thr Pro Arg Trp Glu Lys His
```

Asp	Gln	Gly	Gly	Ser	Lys	Lys	Ala	Lys	Lys	Thr	Gln	Ala	Ser	Gly	Gly
225					230					235					240
Glu	Asn	Gln	Lys	Gln	Leu	Ser	His	Glu	He	Gln	Val	Gly	Trp	Gly	Asn
				245					250					255	
Lys	Gly	Leu	Arg	Arg	Asp	Glu	Asp	Ala	Lys	Glu	Thr	Gln	He	Ala	Thr
			260					265					270		
Lys	Lys	Lys	Leu	Arg	Glu	He	Arg	Glu	Lys	Asp	Trp	Val	Val	He	Gln
		275					280					285			
Ala	Leu	Trp	Trp	Gly	Asn	Arg	Arg	Gln	Val	Ala	Ser	Glu	lle	Tyr	Arg
	290					295					300				
Glu	Phe	Glu	He	Leu	Cys	Trp	Glu	Asn	Gln	Asn	Trp	Ile	Gly	Gly	Glu
305					310					315					320
His	Arg	Ala	Glu	He	Gln	Ala	Ser	Glu	Lys	Arg	Asp	Gln	Arg	Lys	Asp
				325					330					335	
Gly	Cys	Glu	Asp	Gly	Thr	Asn	lle	Leu	Ala	Pro	Glu	Ala	Glu	He	Gln
			340					345					350		
Glu	G]n		Lys	Gly	Glu	Thr		Val	Glu	Thr	Gln		Asn	Glu	Pro
		355					360					365			
Leu		Glu	Glu	Asp	Gly		Asp	Ile	Gln	Ser		Gly	Arg	Arg	Glu
	370					375		<u>_</u> .			380				
	Lys	Gly	Glu	Asp		Lys	Asp	Thr	Gln		Leu	Gly	Arg	Lys	
385					390			_		395					400
GIn	G1 y	GIn	Leu		Asn	Glu	Phe	Ser		Lys	He	His	He		Lys
0.1			61	405				61	410		C.1			415	6.1
GTy	Lys	Asn		Glu	His	11e	Arg	Gly	Glu	Asp	ыу	Ala		lhr	GIn
7.1	C	C1	420	C1		т	C1	425	1	TI.	C	C1	430	Α	C1
11e	ser		Ser	GIY	Asn	ırp		Lys	Leu	Inr	ser		116	Asp	GIŸ
C1	Mat	435	Con	A 1 .s.	C1	Т	440	Luc	Aan	Cla	Cla	445	C1	Clu	C1
GIU		nis	5er	ATA	GIU	-	Lys	Lys	ASP	GIII	460	116	GIY	GIY	GIU
Aan	450	Ma	C1	110	Cla	455	Cl _n	G1 y	Luc	Ana		Lou	Ana	C1u	Vol
465	Gry	MIa	Olu	116	470	116	GIII	GIY	LyS	475	ASII	i.eu	Mg	Gru	480
	Cly	C1u	Acn	Cly		Luc	The	Tro	Ala		Clv	Lve	Glo	Thr	
O I y	Oly	OIU	nsp	485	val	LyS	1111	Trp	490	110	ory	rys	010	495	0111
Ser	Gla	Phe	Ara		Acn	Lev	G1v	Arg		مال	ില	ىرم ا	Ser		Tro
501	0111	, 110	500	001	пар	1, CU	GIY	505	E) 3	.10	,.cu	u	510	o i u	p

Lys Ser Gln Lys Gln Met Gly Ser Glu Asn Gly Thr Glu Ile Gln Ala Pro Val Glu Arg Asn Gln Arg Glu Pro Gly Gly Glu Asp Gly Val Lys Thr Gln Arg Pro Lys Arg Glu Asn Glu Asp Gln Leu Asp Ser Glu lle Gly Gly Ser His Ser Pro Gly Arg Arg Asn Trp Glu Leu lle Gly Lys Asp Val Ala Glu Asn Gln Ala Ser Glu Lys Arg Asn Gln Arg Glu Val Gly Asn Glu Asp Gly Arg Met Ile Trp Arg Leu Arg Gly Lys Asn Trp Arg Leu Arg Ala Lys Lys Gln Thr Val Lys Lys

<210> 4913

<211> 164

<212> PRT

<213> Homo sapiens

<400> 536

Met Asn Val Leu Val Trp Glu Asp Cys lle Ala Glu Gln Ala Glu Val Leu His Asn Asp Ser Tyr Gly Val Ile Ile Asp Cys Ser Pro Lys Gly Met Phe Ser Leu Asn Cys Thr Ser Gln Ser Ala Cys His Gly His Thr Met Phe Ser Trp Ser Glu Gln Asn Gly Gln Met Val Glu Met lle Arg Ser Met Ala Arg Val Pro Ile Ile Trp Lys His Gly Gly Ile Val Ala Pro Gln Pro Gln Met 11e Trp Pro Ala Val Gly Ala Lys His Lys Asp Leu Trp Lys Leu Leu Met Ala Leu Asn Lys Ile Lys Ile Trp Glu Arg

 11e
 Lys
 Lys
 His
 Leu
 Glu
 Gly
 His
 Ser
 Arg
 Asn
 Leu
 Asp
 11e
 Ala
 Lys

 Leu
 Lys
 Glu
 Gln
 11e
 Phe
 Lys
 Ala
 Ser
 Gln
 Ala
 His
 Leu
 Thr
 Leu
 Met

 130
 135
 135
 140
 140
 11e
 Ala
 A

<210> 4914

<211> 236

<212> PRT

<213> Homo sapiens

<400> 537

Met Ser Ser Thr Glu Ser Ala Gly Arg Thr Ala Asp Lys Ser Pro Arg

1 5 10 15

Gln Gln Val Asp Arg Leu Leu Val Gly Leu Arg Trp Arg Arg Leu Glu

Gln Gln Val Asp Arg Leu Leu Val Gly Leu Arg Trp Arg Arg Leu Glu 20 25 30

Glu Pro Leu Gly Phe Ile Lys Val Leu Gln Trp Leu Phe Ala Ile Phe 35 40 45

Ala Phe Gly Ser Cys Gly Ser Tyr Ser Gly Glu Thr Gly Ala Met Val
50 55 60

Arg Cys Asn Asn Glu Ala Lys Asp Val Ser Ser lle IIe Val Ala Phe 65 70 75 80

Gly Tyr Pro Phe Arg Leu His Arg Ile Gln Tyr Glu Met Pro Leu Cys
85 90 95

Asp Glu Gly Ser Ser Ser Lys Thr Met His Leu Met Gly Asp Phe Ser 100 105 110

Ala Pro Ala Glu Phe Phe Val Thr Leu Gly Ile Phe Ser Phe Phe Tyr
115
120
125

Thr Met Ala Ala Leu Val Ile Tyr Leu Arg Phe His Asn Leu Tyr Thr 130 135 140

Glu Asn Lys Arg Phe Pro Leu Val Asp Phe Cys Val Thr Val Ser Phe 145 150 155 160 Thr Phe Phe Trp Leu Val Ala Ala Ala Ala Trp Gly Lys Gly Leu Thr Asp Val Lys Gly Ala Thr Arg Pro Ser Ser Leu Thr Ala Ala Met Ser Val Cys His Gly Glu Glu Ala Val Cys Ser Ala Gly Ala Thr Pro Ser Met Gly Leu Ala Asn Ile Ser Val Val Arg Pro Val Ala Thr Ala Gly Ser Ser Thr Ser Pro Ala Ala Gln Ala Cys Pro Ser

<210> 4915

<211> 390

<212> PRT

<213> Homo sapiens

<400> 538

Met Cys Ile Cys His Leu Pro Cys Arg Pro Val Lys Pro Asn lle Ile Gly Glu Gln Ile Thr Ser Lys Met Gly Ala His Tyr His Cys Ile Ile Cys Ser Ala Thr Ile Thr Arg Arg Thr Asp Met Leu Gly His Val Arg Arg His Met Asn Lys Gly Glu Thr Lys Ser Ser Tyr Ile Ala Ala Ser Thr Ala Lys Pro Pro Lys Glu lle Leu Lys Glu Ala Asp Thr Asp Val Gln Val Cys Pro Asn Tyr Ser Ile Pro Gln Lys Thr Asp Ser Tyr Phe Asn Pro Lys Met Lys Leu Asn Arg Gln Leu lle Phe Cys Thr Leu Ala Ala Leu Ala Glu Glu Arg Lys Pro Leu Glu Cys Leu Asp Ala Phe Gly

Ala Thr Gly Ile Met Gly Leu Gln Trp Ala Lys His Leu Gly Asn Ala

Val	Lys	Val	Thr	He	Asn	Asp	Leu	Asn	Glu	Asn	Ser	Val	Thr	Leu	lle
145					150					155					160
Gln	Glu	Ser	Cys	His	Leu	Asn	Lys	Leu	Lys	Val	Val	Val	Asp	Ser	Lys
				165					170					175	
Glu	Lys	Glu	Lys	Ser	Asp	Asp	11e	Leu	Glu	Glu	Gly	Glu	Lys	Asn	Leu
			180					185					190		
Gly	Asn	lle	Lys	Val	Thr	Lys	Met	Asp	Ala	Asn	Val	Leu	Met	His	Leu
		195					200					205			
Arg	Ser	Phe	Asp	Phe	Ile	His	Leu	Asp	Pro	Phe	Gly	Thr	Ser	Val	Asn
	210					215					220				
Tyr	Leu	Asp	Ser	Ala	Phe	Arg	Asn	lle	Arg	Asn	Leu	Gly	Ile	Val	Ser
225					230					235					240
Val	Thr	Ser	Thr	Asp	lle	Ser	Ser	Leu	Tyr	Ala	Lys	Ala	Gln	His	Val
				245					250					255	
Ala	Arg	Arg	His	Tyr	Gly	Cys	Asn	Île	Val	Arg	Thr	Glu	Tyr	Tyr	Lys
			260					265					270		
Glu	Leu	Ala	Ala	Arg	lle	Val	Val	Ala	Ala	Val	Ala	Arg	Ala	Ala	Ala
		275					280					285			
Arg	Cys	Asn	Lys	Gly	Ile	Glu	Val	Leu	Phe	Ala	Val	Ala	Leu	Glu	His
	290					295					300				
Phe	Val	Leu	Val	Va]	Val	Arg	Val	Leu	Arg	Gly	Pro	Thr	Ser	Ala	Asp
305					310					315					320
Glu	Thr	Ala	Lys	Lys	He	Gln	Tyr	Leu	lle	His	Cys	Gln	Trp	Cys	Glu
				325					330					335	
Glu	Arg	He	Phe	Gln	Lys	Asp	Gly	Asn	Met	Val	Glu	Asp	Tyr	Ser	Ala
			340					345					350		
Asn	Phe	Val	Ile	Ser	Tyr	Thr	Gly	Phe	Pro	Phe	Va]	Asn	Arg	Gln	Asp
		355					360					365			
He	Arg	Lys	Thr	His	He	Asp	Ser	Cys	Leu	Val	Thr	Va]	Met	Glu	Ala
	370					375					380				
Cys	Leu	Glu	Arg	Gln	Gln										
385					390										

<210> 4916

<211> 619

<212> PRT <213> Homo sapiens <400> 539 Met Ala Arg Leu Gln Arg Arg Ala Ser Gln Arg Arg Gln Gly Gly Thr Trp Gly Leu Arg Val Val Gln Glu Pro Gly Gly His Leu Tyr lle Trp Leu Ala Ser Glu Lys Ala His Glu Arg Gln Arg Ala Val His Ser Cys Met Ile Leu Leu Lys Phe Leu Asn His Asn Gly Tyr Leu Asp Pro Lys Glu Asp Phe Lys Arg Ile Gly Gln Leu Val Gly Ile Leu Gly Met Leu Cys Gln Asp Pro Asp Arg Ala Thr Gln Arg Cys Ser Leu Glu Gly Ala Ser His Leu Tyr Gln Leu Leu Met Cys His Lys Arg Glu Ala Leu Gln Ala Glu Ser Gln Ala Pro Lys Glu Leu Ser Gln Ala His Ser Asp Gly Ala Pro Leu Trp Asn Ser Arg Asp Gln Lys Ala Thr Pro Leu Gly Pro Gln Glu Met Ala Lys Asn His 11e Phe Gln Leu Cys Ser Phe Gln Val lle Lys Asp lle Met Gln Gln Leu Thr Leu Ala Glu Leu Ser Asp Leu lle Trp Thr Ala lle Asp Gly Leu Gly Ser Thr Ser Pro Phe Arg Val Gln Ala Ala Ser Glu Met Leu Leu Thr Ala Val Gln Glu His Gly Ala Lys Leu Glu Ile Val Ser Ser Met Ala Gln Ala Ile Arg Leu Arg Leu Cys Ser Val His Ile Pro Gln Ala Lys Glu Lys Thr Leu His Ala Ile

Thr Leu Leu Ala Arg Ser His Thr Cys Glu Leu Val Ala Thr Phe Leu

Asn	He	Ser	Ile	Pro	Leu	Asp	Ser	His	Thr	Phe	Gln	Leu	Trp	Arg	Ala
			260					265					270		
Leu	Gly	Ala	Glu	Gln	Pro	Thr	Ser	His	Leu	Val	Leu	Thr	Thr	Leu	Leu
		275					280					285			
Ala	Cys	Leu	Gln	Glu	Arg	Pro	Leu	Pro	Thr	Gly	Ala	Ser	Asp	Ser	Ser
	290					295					300				
Pro	Cys	Pro	Lys	Glu	Lys	Thr	Tyr	Leu	Arg	Leu	Leu	Ala	Ala	Met	Asn
305					310					315					320
Met	Leu	His	Glu	Leu	Gln	Phe	Ala	Arg	Glu	Phe	Lys	Gln	Ala	Val	Gln
				325					330					335	
Glu	Gly	Tyr	Pro	Lys	Leu	Phe	Leu	Ala	Leu	Leu	Thr	Gln	Met	His	Tyr
			340					345					350		
Val	Leu	Glu	Leu	Asn	Leu	Pro	Ser	Glu	Pro	Gln	Pro		Gln	G1n	Ala
		355					360					365			
Gln		Ala	Ala	Val	Pro	Ser	Pro	Gln	Ser	Cys		Thr	Ser	Leu	Glu
	370		_			375					380				
	Leu	Lys	Ser	Leu		Ser	Thr	Thr	Gly		Trp	His	Asp	Phe	
385		61	,	01	390	C	T.	61	,	395	T)	T)	7.1		400
HIS	Leu	Glu	Leu		Gly	Ser	Irp	Glu		Phe	Ihr	lhr	He		Ihr
Tum	Dno	1	Cly	405	Clu	Lou	Lou	A1 o	410	110	Mot	Vo.1	C1p	415	Шi.a
lyr	Pro	Lys	420	vai	Gly	Leu	Leu	425	Arg	мта	мет	vai	430	ASII	піѕ
Cvc	Ara	Gln		Pro	A1a	Val	Lou		Gln	Lou	Lou	Pro		Lou	Cl _D
Cys	ni g	435	116	110	MIG	vai	440	Mg	0111	Leu	Leu	445	561	Leu	OIII
Ser	Pro		Glu	Arø	Glu	Arg		Val	Ala	He	l eu		Leu	Thr	Lvs
001	450	0111	0.0	6	0.0	455	2,0				460		200		25,0
Phe		Tvr	Ser	Pro	Val	Leu	Leu	Glu	Val	Leu		Lvs	Gln	Ala	Ala
465		•			470					475		,			480
Leu	Thr	Val	Leu	Ala	Gln	Gly	Leu	His	Asp	Pro	Ser	Pro	Glu	Val	Arg
				485					490					495	
Val	Leu	Ser	Leu	Gln	Gly	Leu	Ser	Asn	Ile	Leu	Phe	His	Pro	Asp	Lys
			500					505					510		
Glu	Arg	Asp	G1 y	He	Arg	Ala	Ala	Ala	Met	Ala	Leu	Phe	Gly	Asp	Leu
		515					520					525			
Val	Ala	Ala	Met	Ala	Asp	Arg	Glu	Leu	Ser	Gly	Leu	Arg	Thr	Gln	Val
	530					535					540				

His Gln Ser Met Val Pro Leu Leu Leu His Leu Lys Asp Gln Cys Pro 555 550 Ala Val Ala Thr Gln Ala Lys Phe Thr Phe Tyr Arg Cys Ala Val Leu 565 570 575 Leu Arg Trp Arg Leu Leu His Thr Leu Phe Cys Thr Leu Ala Trp Glu 590 580 585 Arg Gly Leu Ser Ala Arg His Phe Leu Trp Thr Cys Leu Ala Thr Pro 600 605 Ser Ala Thr Thr Pro Arg Pro Cys Ser Arg Cys 610 615

<210> 4917

<211> 1103

<212> PRT

<213> Homo sapiens

<400> 540

Met His Phe Leu Ala Ala Gln Asn Gln Phe Leu His Ser Pro Phe Leu

1 5 10 15

Glu Arg Pro Met Asp Met Pro Tyr Met Ile Phe Asp Pro Asn Asn Pro
20 25 30

Leu Met Thr Gly Gln Leu Leu Gly Ser Ser Leu Thr Gln Met Pro Pro
35 40 45

Gln Ala Ser Ser Ser His Thr Thr Ala Pro Thr Thr Val Ala Ala Ser 50 55 60

Leu Lys Arg Lys Leu Asp Asp Lys Glu Asp Asn Asn Cys Ser Glu Lys
65 70 75 80

Glu Gly Gly Asn Ser Gly Glu Asp Gln His Arg Asp Lys Arg Leu Arg

85 90 95

Thr Thr Ile Thr Pro Glu Gln Leu Glu Ile Leu Tyr Glu Lys Tyr Leu 100 105 110

Leu Asp Ser Asn Pro Thr Arg Lys Met Leu Asp His 11e Ala Arg Glu 115 120 125

Val Gly Leu Lys Lys Arg Val Val Gln Val Trp Phe Gln Asn Thr Arg

	130					135					140				
Ala	Arg	Glu	Arg	Lys	Gly	Gln	Phe	Arg	Ala	Val	Gly	Pro	Ala	Gln	Ser
145					150					155					160
His	Lys	Arg	Cys	Pro	Phe	Cys	Arg	Ala	Leu	Phe	Lys	Ala	Lys	Ser	Ala
				165					170					175	
Leu	Glu	Ser	His	He	۸rg	Ser	Arg	His	Trp	Asn	Glu	Gly	Lys	Gln	Ala
			180					185					190		
Gly	Tyr	Ser	Leu	Pro	Pro	Ser	Pro	Leu	He	Ser	Thr	Glu	Asp	Gly	Gly
		195					200					205			
Glu	Ser	Pro	Gln	Lys	Tyr	lle	Tyr	Phe	Asp	Tyr	Pro	Ser	Leu	Pro	Leu
	210					215					220				
Thr	Lys	Ile	Asp	Leu	Ser	Ser	Glu	Asn	Glu	Leu	Ala	Ser	Thr	Val	Ser
225					230					235					240
Thr	Pro	Val	Ser	Lys	Thr	Ala	Glu	Leu	Ser	Pro	Lys	Asn	l.eu	Leu	Ser
				245					250					255	
Pro	Ser	Ser	Phe	Lys	Ala	Glu	Cys	Ser	Glu	Asp	Val	Glu	Asn	Leu	Asn
			260					265					270		
Ala	Pro	Pro	Ala	Glu	Ala	Gly	Tyr	Asp	Gln	Asn	Lys	Thr	Asp	Phe	Asp
		275					280					285			
Glu	Thr	Ser	Ser	lle	Asn	Thr	Ala	lle	Ser	Asp	Ala	Thr	Thr	Gly	Asp
	290					295					300				
Glu	Gly	Asn	Thr	Glu	Met	Glu	Ser	Thr	Thr	Gly	Ser	Ser	Gly	Asp	Val
305					310					315					320
Lys	Pro	Ala	Leu	Ser	Pro	Lys	Glu	Pro	Lys	Thr	Leu	Asp	Thr	Leu	Pro
				325					330					335	
Lys	Pro	Ala	Thr	Thr	Pro	Thr	Thr				Asp	Asp	Lys	Phe	Leu
			340										350		
Phe	Ser		Thr	Ser	Pro	Ser	He	His	Phe	Asn	Asp		Asp	Gly	Asp
		355					360					365			
His		Gln	Ser	Phe	Tyr		Thr	Asp	Asp	Pro		Asp	Asn	Ala	Asp
	370					375					380				
-	Ser	Glu	Thr	Ser		He	Ala	Asp	Pro		Ser	Pro	Asn	Pro	
385	~				390		_		~	395			ъ	0.3	400
Gly	Ser	Ser	Asn		Phe	Lys	Ser	Lys		Asn	Asp	Arg	Pro		His
		n.		405					410	0.			., .	415	,
Lys	Arg	Phe	Arg	Thr	GIn	Met	Ser	Asn	Leu	GIn	Leu	Lys	Val	Leu	Lys

			420					425					430		
Ala	Cys	Phe	Ser	Asp	Tyr	Arg	Thr	Pro	Thr	Met	G1n	Glu	Cys	Glu	Met
		435					440					445			
Leu	Gly	Asn	Glu	Пе	Gly	Leu	Pro	Lys	Arg	Val	Val	G1n	Val	Trp	Phe
	450					455					460				
Gln	Asn	Ala	Arg	Ala	Lys	Glu	Lys	Lys	Phe	Lys	He	Asn	He	Gly	Lys
465					470					475					480
Pro	Phe	Met	Пе	Asn	Gln	Gly	Gly	Thr	Glu	Gly	Thr	Lys	Pro	Glu	Cys
				485					490					495	
Thr	Leu	Cys	Gly	Val	Lys	Tyr	Ser	Ala	Arg	Leu	Ser	Ile	Arg	Asp	His
			500					505					510		
Tle	Phe	Ser	Lys	Gln	His	He	Ser	Lys	Val	Arg	Glu	Thr	Val	Gly	Ser
		515					520					525			
Gln	Leu	Asp	Arg	Glu	Lys	Asp	Tyr	Leu	Λla	Pro	Thr	Thr	Val	Arg	Gln
	530					535					540				
Leu	Met	Ala	Gln	Gln	Glu	Leu	Asp	Arg	Ile	Lys	Lys	Ala	Ser	Asp	Val
545					550					555					560
Leu	Gly	Leu	Thr	Val	Gln	Gln	Pro	Gly	Met	Met	Asp	Ser	Ser	Ser	Leu
				565					570					575	
His	Gly	He	Ser	Leu	Pro	Thr	Ala	Tyr	Pro	Gly	Leu	Pro	Gly	Leu	Pro
			580					585					590		
Pro	Val	Leu	Leu	Pro	Gly	Met	Asn	Gly	Pro	Ser	Ser	Leu	Pro	Gly	.Phe
		595					600					605			
Pro	Gln	Asn	Ser	Asn	Thr	Leu	Thr	Pro	Pro	Gly	Ala	Gly	Met	Leu	Gly
	610					615					620				
Phe	Pro	Thr	Ser	Ala	Thr	Ser	Ser	Pro	Ala	Leu	Ser	Leu	Ser	Ser	Ala
625					630					635					640
Pro	Thr	Lys	Pro	Leu	Leu	Gln	Thr	Pro							
				645					650					655	
Pro	Pro	Pro	Pro	Ser	Ser	Ser	Leu	Ser	Gly	Gln	Gln	Thr		Gln	Gln
			660					665					670		
Asn	Lys		Ser	Glu	Lys	Lys		Thr	Lys	Pro	Asn	Lys	Val	Lys	Lys
		675					680				_@	685		_	
He		Glu	Glu	Glu	Leu		Ala	Thr	Lys	Pro		Lys	His	Pro	Lys
	690				_	695		_			700		_		
Lys	Glu	Glu	Lys	He	Ser	Ser	Ala	Leu	Ser	Va]	Leu	Gly	Lys	Val	Val

705					710					715					720
G1 y	Glu	Thr	His	Val	Asp	Pro	Пe	Gln	Leu	Gln	Ala	Leu	Gln	Asn	Ala
				725					730					735	
lle	Ala	Gly	Asp	Pro	Ala	Ser	Phe	He	Gly	G1 y	Gln	Phe	Leu	Pro	Tyr
			740					745					750		
Phe	He	Pro	G1 y	Phe	Ala	Ser	Tyr	Phe	Thr	Pro	Gln	Leu	Pro	Gly	Thr
		755					760					765			
Val	Gln	Gly	Gly	Tyr	Phe	Pro	Pro	Val	Cys	Gly	Met	Glu	Ser	Leu	Phe
	770					775					780				
Pro	Tyr	Gly	Pro	Thr	Met	Pro	Gln	Thr	Leu	Ala	Gly	Leu	Ser	Pro	Gly
785					790					795					800
Ala	Leu	Leu	Gln	Gln	Tyr	Gln	Gln	Tyr	Gln	Gln	Asn	Leu	Gln	Glu	Ser
				805					810					815	
Leu	Gln	Lys	Gln	Gln	Lys	Gln	Gln	Gln	Glu	Gln	Gln	Gln	Lys	Pro	Ala
			820					825					830		
Gln	Ala	Lys	Thr	Ser	Lys	Val	Glu	Ser	Asp	Gln	Pro	Gln	Asn	Ser	Asn
		835					840					845			
Asp	Ala	Ser	Glu	Thr	Lys	Glu	Asp	Lys	Ser	Thr	Ala	Thr	Glu	Ser	Thr
	850					855					860				
Lys	Glu	Glu	Pro	Gln	Leu	Glu	Ser	Lys	Ser	Ala	Asp	Phe	Ser	Asp	Thr
865					870					875					880
Tyr	Val	Val	Pro	Phe	Val	Lys	Tyr	Glu	Phe	He	Cys	Arg	Lys	Cys	Gln
				885					890					895	
Met	Met	Phe	Thr	Asp	Glu	Asp	Ala	Ala	Val	Asn	His	Gln	Lys	Ser	Phe
			900					905					910		
Cys	Tyr	Phe	Gly	Gln	Pro	Leu	Île	Asp	Pro	Gln	Glu	Thr	Val	Leu	Arg
		915					920					925			
Val	Pro	Val	Ser	Lys	Tyr	Gln	Cys	Leu	Ala	Cys	Asp	Val	Ala	lle	Ser
	930					935					940				
Gly	Asn	Glu	Ala	Leu	Ser	Gln	His	Leu	Gln	Ser	Ser	Leu	His	Lys	Glu
945					950					955					960
Lys	Thr	lle	Lys	Gln	Ala	Met	Arg	Asn	Ala	Lys	Glu	His	Val	Arg	Leu
				965					970					975	
Leu	Pro	His	Ser	Val	Cys	Ser	Pro	Asn	Pro	Asn	Thr	Thr	Ser	Thr	Ser
			980					985					990		
Gln	Ser	Ala	Ala	Ser	Ser	Asn	Asp	Thr	Tyr	Pro	Hic	Leu	Ser	Cvs	Phe

1000 1005 Ser Met Lys Ser Trp Pro Asn Ile Leu Phe Gln Ala Ser Ala Arg Arg 1015 Ala Ala Ser Pro Pro Ser Ser Pro Pro Ser Leu Ser Leu Pro Ser Thr 1025 1030 1035 1040 Val Thr Ser Ser Leu Cys Ser Thr Ser Gly Val Gln Thr Ser Leu Pro 1045 1050 Thr Glu Ser Cys Ser Asp Glu Ser Asp Ser Glu Leu Ser Gln Lys Leu 1065 Glu Asp Leu Asp Asn Ser Leu Glu Val Lys Ala Lys Pro Ala Ser Gly 1075 1080 1085 Leu Asp Gly Asn Phe Asn Ser Ile Arg Met Asp Met Phe Ser Val 1090 1095 1100 <210> 4918 <211> 727 <212> PRT <213> Homo sapiens <400> 541 Met Gln Gln Lys Thr Lys Leu Phe Pro Gln Ala Leu Lys Tyr Ser lle 10

Net Gin Gin Lys Inr Lys Leu Phe Pro Gin Ala Leu Lys Tyr Ser Ile

1 5 10 15

Pro His Leu Gly Lys Cys Met Gln Lys Gln His Leu Asn His Tyr Asn
20 25 30

Phe Ala Asp His Cys Tyr Asn Arg Ile Lys Leu Lys Lys Tyr His Leu

Val	Glu		Lys	His	Gly	Gly		Val	Tyr	Tyr	Gln		Gly	Cys	Cys
		115					120					125			
Leu	Val	Arg	Ser	Lys	Asp		Glu	Ala	Asp	Asn	Asp	Asn	Tyr	Glu	Val
	130					135					140				
Leu	Phe	Λsn	Leu	Glu	Glu	Leu	Lys	Leu	Asp	Gln	Pro	Phe	He	Asp	Cys
145					150					155					160
Пe	Arg	Val	Ala	Pro	Asp	G]u	Lys	Tyr	Val	Ala	Ala	Lys	lle	Arg	Thr
				165					170					175	
Glu	Asp	Ser	Glu	Ala	Ser	Thr	Cys	Val	He	He	Lys	Leu	Ser	Asp	Gln
			180					185					190		
Pro	Val	Met	Glu	Ala	Ser	Phe	Pro	Asn	Val	Ser	Ser	Phe	Glu	Trp	Val
		195					200					205			
Lys	Λsp	Glu	Glu	Asp	Glu	Asp	Val	Leu	Phe	Tyr	Thr	Phe	Gln	Arg	Asn
	210					215					220				
Leu	Arg	Cys	His	Asp	Val	Tyr	Arg	Ala	Thr	Phe	Gly	Asp	Asn	Lys	Arg
225					230					235					240
Asn	Glu	Arg	Phe	Tyr	Thr	Glu	Lys	Asp	Pro	Ser	Tyr	Phe	Val	Phe	Leu
				245					250					255	
Tyr	Leu	Thr	Lys	Asp	Ser	Arg	Phe	Leu	Thr	lle	Asn	Ile	Met	Asn	Lys
			260					265					270		
Thr	Thr	Ser	Glu	Val	Trp	Leu	lle	Asp	Gly	Leu	Ser	Pro	Trp	Asp	Pro
		275					280					285			
Pro	Val	Leu	lle	Gln	Lys	Arg	lle	His	Gly	He	Leu	Tyr	Tyr	Val	Glu
	290					295					300				
His	Arg	Asp	Asp	Glu	Leu	Tyr	IJe	Leu	Thr	Asn	Val	Gly	Glu	Pro	Thr
305					310					315					320
Glu	Phe	Lys	Leu	Met	Λrg	Thr	Ala	Ala	Asp	Thr	Pro	Ala	Ile	Met	Asn
				325					330					335	
Trp	Asp	l.eu	Phe	Phe	Thr	Met	Lys	Arg	Asn	Thr	Lys	Val	Ile	Asp	Leu
			340					345					350		
Asp	Met	Phe	Lys	Asp	His	Cys	Val	Leu	Phe	Leu	Lys	His	Ser	Asn	Leu
		355					360					365			
Leu	Tyr	Va]	Asn	Va]	lle	Gly	Leu	Ala	Asp	Asp	Ser	Val	Arg	Ser	Leu
	370					375					380				
Lys	Leu	Pro	Pro	Trp	Ala	Cys	Gly	Phe	He	Met	Asp	Thr	Asn	Ser	Asp
385					390					395					400

Pro	Lys	Asn	Cys	Pro	Phe	Gln	Leu	Cys	Ser	Pro	He	Arg	Pro	Pro	Lys
				405					410					415	
Tyr	Tyr	Thr	Tyr	Lys	Phe	Ala	Glu	Gly	Lys	Leu	Phe	Glu	Glu	Thr	Gly
			420					425					430		
His	Glu	Asp	Pro	lle	Thr	Lys	Thr	Ser	Arg	Val	Leu	Arg	Leu	Glu	Ala
		435					440					445			
Lys	Ser	Lys	Asp	Gly	Lys	Leu	Val	Pro	Met	Thr	Val	Phe	His	Lys	Thr
	450					455					460				
Asp	Ser	Glu	Asp	Leu	Gln	Lys	Lys	Pro	Leu	Leu	Val	His	Val	Tyr	Gly
465					470					475					480
Ala	Tyr	Gly	Met	Asp	Leu	Lys	Met	Asn	Phe	Arg	Pro	Glu	Arg	Arg	Va]
				485					490					495	
Leu	Val	Asp	Asp	Gly	Trp	lle	Leu	Ala	Tyr	Cys	His	Val	Arg	Gly	Gly
			500					505					510		
G1 y	Glu	Leu	Gly	Leu	Gln	Trp	His	Ala	Asp	Gly	Arg	Leu	Thr	Lys	Lys
		515					520					525			
Leu	Asn	Gly	Leu	Ala	Asp	Leu	Glu	Ala	Cys	Ile	Lys	Thr	Leu	His	Gly
	530					535					540				
Gln	Gly	Phe	Ser	Gln	Pro	Ser	Leu	Thr	Thr	Leu	Thr	Ala	Phe	Ser	Ala
545					550					555					560
Gly	Gly	Val	Leu	Ala	Gly	Ala	Leu	Cys	Asn	Ser	Asn	Pro	Glu	Leu	Val
				565					570					575	
Arg	Ala	Va]	Thr	Leu	Glu	Ala	Pro	Phe	Leu	Asp	Val	Leu	Asn	Thr	Met
			580					585					590		
Met	Asp	Thr	Thr	Leu	Pro	Leu	Thr	Leu	Glu	Glu	Leu	Glu	Glu	Trp	Gly
		595					600					605			
Asn	Pro	Ser	Ser	Asp	Glu	Lys	His	Lys	Asn	Tyr	Ile	Lys	Arg	Tyr	Cys
	610					615					620				
Pro	Tyr	Gln	Asn	He	Lys	Pro	Gln	His	Tyr	Pro	Ser	He	His	He	Thr
625					630					635					640
Ala	Tyr	Glu	Asn	Asp	Glu	Arg	Val	Pro	Leu	Lys	Gly	lle	Val	Ser	Tyr
				645					650					655	
Thr	Glu	Lys	Leu	Lys	Glu	Ala	He	Ala	Glu	His	Ala	Lys	Asp	Thr	Gly
			660					665					670		
Glu	Gly	Tyr	Gln	Thr	Pro	Asn	He	He	Leu	Asp	11e	GIn	Pro	Gly	Gly
		675					680					685			

Asn His Val lle Glu Asp Ser His Lys Lys lle Thr Ala Gln Ile Lys Phe Leu Tyr Glu Glu Leu Gly Leu Asp Ser Thr Ser Val Phe Glu Asp Leu Lys Lys Tyr Leu Lys Phe <210> 4919 <211> 378 <212> PRT <213> Homo sapiens <400> 542 Met Glu Asp Arg Asp Glu Ser Glu Val Ser Asp Glu Gly Gly Ser Pro lle Ser Ser Glu Gly Gln Glu Pro Arg Ala Asp Pro Glu Pro Pro Gly Leu Ala Ala Gly Leu Val Gln Gln Asp Leu Val Phe Glu Val Glu Thr Pro Ala Val Leu Pro Glu Pro Val Pro Gln Glu Asp Gly Val Asp Leu Leu Gly Leu His Ser Glu Val Gly Ala Gly Pro Ala Val Pro Pro Gln Ala Cys Lys Ala Pro Ser Ser Asn Thr Asp Leu Leu Ser Cys Leu Leu Gly Pro Pro Glu Ala Ala Ser Gln Gly Pro Pro Glu Asp Leu Leu Ser Glu Asp Pro Leu Leu Leu Ala Ser Pro Ala Pro Pro Leu Ser Val Gln Ser Thr Pro Arg Gly Gly Pro Pro Ala Ala Ala Asp Pro Phe Gly Pro Leu Leu Pro Ser Ser Gly Asn Asn Ser Gln Pro Cys Ser Asn Pro Asp

Leu Phe Gly Glu Phe Leu Asn Ser Asp Ser Val Thr Val Pro Pro Ser

Phe	Pro	Ser	Ala	His	Ser	Ala	Pro	Pro	Pro	Ser	Cys	Ser	Ala	Asp	Phe
			180					185					190		
Leu	His	Leu	Gly	Asp	Leu	Pro	Gly	Glu	Pro	Ser	Lys	Met	Thr	Ala	Ser
		195					200					205	•		
Ser	Ser	Asn	Pro	Asp	Leu	Leu	Gly	Gly	Trp	Ala	Ala	Trp	Thr	Glu	Thr
	210					215					220				
Ala	Ala	Ser	Ala	Val	Λla	Pro	Thr	Pro	Ala	Thr	Glu	Gly	Ser	Pro	Ala
225					230					235					240
Gly	Phe	Pro	Pro	Gly	Gly	Phe	lle	Pro	Lys	Thr	Ala	Thr	Thr	Pro	Lys
				245					250					255	
Gly	Ser	Ser	Ser	Trp	Gln	Thr	Ser	Arg	Pro	Pro	Ala	Gln	Gly	Ala	Ser
			260					265					270		
Trp	Pro	Pro	Gln	Ala	Lys	Pro	Pro	Pro	Lys	Ala	Cys	Thr	Gln	Pro	Arg
		275					280					285			
Pro	Asn	Tyr	Ala	Ser	Asn	Phe	Ser	Val	lle	G1 y	Ala	Arg	Glu	Glu	Arg
	290					295					300				
Gly	Val	Arg	Ala	Pro	Ser	Phe	Gly	Glu	Ser	Pro	Thr	Leu	Cys	Cys	Cys
305					310					315					320
Gly	Val	Cys	Arg	Asp	Leu	Pro	Glu	Leu	Ala	Lys	Gly	Arg	Thr	Tyr	Val
				325					330					335	
Val	Phe	Phe	lle	Ser	Glu	Pro	Ser	Pro	G1 y	Thr	Gly	Thr	Ala	Lys	Pro
			340					345					350		
Leu	Arg	Glu	Arg	Arg	Gly	Phe	Arg	Cys	Val	Phe	Thr	Gly	Leu	Ser	Leu
		355					360					365			
Gly	Gln	Ala	Ser	Gly	Ser	Leu	Leu	Phe	Gln						
	370					375									

<211> 168

<212> PRT

<213> Homo sapiens

<400> 543

Met Glu Glu Ile Arg Phe Pro Arg Thr Leu Gly Pro Glu Ala Lys Ser 1 5 10 15 Leu Leu Ser Gly Leu Leu Lys Lys Asp Pro Lys Gln Arg Leu Gly Gly 25 Gly Ser Glu Asp Ala Lys Glu Ile Met Gln His Arg Phe Phe Ala Gly 45 40 lle Val Trp Gln His Val Tyr Glu Lys Lys Val Arg Leu Leu Pro Ala 60 55 Tyr Ser Arg Ala Arg Met Leu Pro Thr Tyr Pro His Ser Arg Met His 70 75 Val Ala Arg Ser Pro Asp Phe Pro His Thr Arg Pro His Leu Arg Ser 90 95 85 Leu Leu Gln Ser Trp Tyr Lys Glu Gly Leu Ala Ala Pro Thr Ser Ala 100 105 Pro Gly Ala Gln Arg Leu Trp His Cys Arg Val Pro Pro Gly Asn Trp 125 120 Pro Gly Pro His Phe Leu Leu Pro Ser Glu Val Cys His Thr Leu Ser 140 130 135 Phe Leu Leu Pro Ser Glu Val Cys His Thr Leu Ser Ala Ser Leu Gly 150 160 155 Val Pro Phe Pro Asp Ala Val Gln 165

<210> 4921

<211> 219

<212> PRT

<213> Homo sapiens

<400> 544

Met Val Gly Gly Arg Ser Cys Pro Leu Arg Gln Val Trp Val Gly Asn
1 5 10 15

Lys Ser Gly Leu Leu Val Gly Ala Glu Ala Gly Gly Ser Ala Ala Asp 20 25 30

Gly Val Thr Pro Pro Gln Glu Cys lle Leu Ser Gly lle Met Ser Val 35 40 45

Asn Gly Lys Lys Val Leu His Met Asp Arg Asn Pro Tyr Tyr Gly Gly
50 55 60

Glu Ser Ser Ser Ile Thr Pro Leu Glu Glu Leu Tyr Lys Arg Phe Gln 70 . Leu Leu Glu Gly Pro Pro Glu Ser Met Gly Arg Gly Arg Asp Trp Asn 85 90 Val Asp Leu Ile Pro Lys Phe Leu Met Ala Asn Gly Gln Leu Val Lys 105 100 Met Leu Leu Tyr Thr Glu Val Thr Arg Tyr Leu Asp Phe Lys Val Val 120 125 Glu Gly Ser Phe Val Tyr Lys Gly Gly Lys Ile Tyr Arg Val Pro Ser 135 140 130 Thr Glu Thr Glu Ala Leu Ala Ser Asn Leu Met Gly Met Phe Glu Lys 150 155 Arg Arg Phe Arg Lys Phe Leu Val Phe Val Ala Asn Phe Asp Glu Asn 165 170 Asp Pro Lys Thr Phe Glu Gly Val Asp Pro Gln Thr Thr Ser Met Arg 180 185 190 Asp Val Tyr Arg Lys Phe Asp Leu Gly Gln Asp Val Ile Asp Phe Thr 200 205 Gly His Ala Leu Ala Leu Tyr Arg Thr Asp Glu 210 215

<210> 4922

<211> 419

<212> PRT

<213> Homo sapiens

<400> 545

 Met Leu Arg Cys
 Phe His Ser Lys
 Gly
 Val
 Asn
 Tyr
 1le
 Asn
 Phe Ser

 1
 5
 10
 15

 Ala Thr Gly
 Lys
 Leu Leu Val
 Ser Val
 Gly
 Val
 Asp
 Pro Glu
 His
 Thr

 20
 25
 30
 30
 30
 11e
 Thr
 Val
 Trp
 Gln
 Glu
 Gly
 Ala
 Lys
 Val
 Ala
 Ser
 Arg
 Gly

 35
 40
 45
 45

 Gly
 His
 Leu
 Glu
 Arg
 Pro
 Asp
 Ser
 Asp

 50
 55
 60
 60

Ihr	GIn	Phe	Val	Ser	Val	Gly	Val	Lys	HIS	Met	Lys	Phe	Irp	Ihr	Leu
65					70					75					80
Ala	Gly	Ser	Ala	Leu	Leu	Tyr	Lys	Lys	Gly	Val	lle	Gly	Ser	Leu	Gly
				85					90					95	
Ala	Ala	Lys	Met	Gln	Thr	Met	Leu	Ser	Val	Ala	Phe	Gly	Ala	Asn	Asn
			100					105					110		
Leu	Thr	Phe	Thr	Gly	Ala	He	Asn	Gly	Asp	Val	Tyr	Val	Trp	Lys	Asp
		115					120					125			
His	Phe	Leu	He	Arg	Leu	Val	Ala	Lys	Ala	His	Thr	Gly	Pro	Val	Phe
	130					135					140				
Thr	Met	Tyr	Thr	Thr	Leu	Arg	Asp	Gly	Leu	He	Val	Thr	G1 y	Gly	Lys
145					150					155					160
Glu	Arg	Pro	Thr	Lys	Glu	Gly	Gly	Ala	Val	Lys	Leu	Trp	Asp	Gln	Glu
				165					170					175	
Met	Lys	Arg	Cys	Arg	Ala	Phe	Gln	Leu	Glu	Thr	Gly	Gln	Leu	Va]	Glu
			180					185					190		
Cys	Val	Arg	Ser	Val	Cys	Arg	Gly	Lys	Gly	Lys	He	Leu	Val	Gly	Thr
		195					200					205			
Lys	Asp	Gly	Glu	Ile	lle	Glu	Val	Gly	Glu	Lys	Asn	Ala	Ala	Ser	Asn
	210					215					220				
He	Leu	11e	Asp	Gly	His	Met	Glu	Gly	Glu	11e	Trp	Gly	Leu	Ala	Thr
225					230					235					240
llis	Pro	Ser	Lys	Asp	Leu	Phe	He	Ser	Ala	Şer	Asn	Asp	Gly	Thr	Ala
				245					250					255	
Arg	Ile	Trp	Asp	Leu	Ala	Asp	Lys	Lys	Leu	Leu	Asn	Lys	Val	Ser	Leu
			260					265					270		
Gly	His	Ala	Ala	Arg	Cys	Ala	Ala	Tyr	Ser	Pro	Asp	Gly	Glu	Met	Val
		275					280					285			
Ala	lle	Gly	Met	Lys	Asn	Gly	Glu	Phe	Va]	lle	Leu	Leu	Val	Asn	Ser
	290					295					300				
Leu	Lys	Val	Trp	Gly	Lys	Lys	Arg	Asp	Arg	Lys	Ser	Ala	lle	Gln	Asp
305					310					315					320
He	Arg	11e	Ser	Pro	Asp	Asn	Arg	Phe	Leu	Ala	Val	Gly	Ser	Ser	Glu
				325					330					335	
His	Thr	Val	Asp	Phe	Tyr	Asp	Leu	Thr	Gln	Gly	Thr	Asn	Leu	Asn	Arg

Ile Gly Tyr Cys Lys Asp Ile Pro Ser Phe Val Ile Gln Met Asp Phe Ser Ala Asp Gly Glu Tyr Ile Gln Met Gln Leu Thr Leu Leu Val Gly Asn His Thr Gln Arg Thr Ala lle Leu Gly Leu Trp Lys lle Ala Arg Lys Trp Met Arg Arg Lys Met Gly Lys Trp Pro Ala Val Val Met Gly Trp Pro

<210> 4923

<211> 699

<212> PRT

<213> Homo sapiens

<400> 546

Met Asn Thr Phe Gln Ala Ser Val Ser Phe Gln Asp Val Thr Val Glu Phe Ser Gln Glu Glu Trp Gln His Met Gly Pro Val Glu Arg Thr Leu Tyr Arg Asp Val Met Leu Glu Asn Tyr Ser His Leu Val Ser Val Gly Tyr Cys Phe Thr Lys Pro Glu Leu Ile Phe Thr Leu Glu Gln Gly Glu Asp Pro Trp Leu Leu Glu Lys Glu Lys Gly Phe Leu Ser Arg Asn Ser Pro Glu Asp Ser Gln Pro Asp Glu Ile Ser Glu Lys Ser Pro Glu Asn Gln Gly Lys His Leu Leu Gln Val Leu Phe Thr Asn Lys Leu Leu Thr Thr Glu Gln Glu Ile Ser Gly Lys Pro His Asn Arg Asp Ile Asn Ile

Phe Arg Ala Arg Met Met Pro Cys Lys Cys Asp Ile Ala Gly Ser Ala

	130					135					140				
Cys	Gln	Gly	Leu	Ser	Leu	Met	Ala	Pro	His	Cys	Gln	Tyr	Ser	Lys	Glu
145					150					155					160
Lys	Лlа	His	G]u	Arg	Asn	Val	Cys	Asp	Lys	Trp	Leu	He	Ser	lle	Lys
				165					170					175	
Asp	Gly	Arg	Thr	Asn	Thr	Gln	Glu	Lys	Ser	Phe	Ala	Tyr	Ser	Lys	lle
			180					185					190		
Val	Lys	Thr	Leu	His	His	Lys	Glu	Glu	Val	Ile	Gln	His	Gln	Thr	Ile
		195					200					205			
Gln	Thr	Leu	Gly	Gln	Asp	Phe	Glu	Tyr	Asn	Glu	Ser	Arg	Lys	Ala	Phe
	210					215					220				
Leu	Glu	Lys	Ala	Ala	Leu	Val	Thr	Ser	Asn	Ser	Thr	His	Pro	Lys	Gly
225					230					235					240
Lys	Ser	Tyr	Asn	Phe	Asn	Lys	Phe	Gly	Glu	Asn	Lys	Tyr	Asp	Lys	Ser
				245					250					255	
Thr	Phe	He	lle	Pro	Gln	Asn	Met	Asn	Pro	Glu	Lys	Ser	His	Tyr	Glu
			260					265					270		
Phe	Asn	Asp	Thr	Gly	Asn	Cys	Phe	Cys	Arg	lle	Thr		Lys	Thr	Leu
		275					280					285			
Thr		Gly	Lys	Ser	Phe		Gln	Lys	Ser	His		Arg	Glu	His	His
	290					295					300				
	Val	His	He	Gly		Lys	Pro	Phe	Glu		Gly	Lys	Ser	Phe	
305			mı		310			0.1		315			m)		320
Arg	Asn	Ser	Thr		Pro	Val	His	GIn		Thr	His	Ala	Thr		Lys
т	С.		т.	325	D.	C	ті.	C1	330	DI	C	т	C1.d	335	ть
lyr	ser	ASP	1 yr	nis	Pro	Cys	ınr	345		rne	ser	iyr	G1 n 350	ser	inr
Dho	Son	Vol	0.10	Cln	Luc	Vol	u; c			110	Lvc	Dro		Clu	Tur
rne	261	355	1112	GIII	rys	vai	360	116	MI g	МІа	Lys	365	1 y 1	010	Tyr
Acn	Glu		Glv	lve	Ser	Cvs		Met	Asn	Ser	Hic		lle	Trn	Pro
изн	370	Cys	Oly	Lys	501	375	501	me c	non	JCI	380	LCu	.1.10	11 p	110
Gln		Ser	His	Thr	Glv		lvs	Pro	Tvr	Glu		Pro	Glu	Cvs	Glv
385	٠,٠٥		,,,,		390	~,u	٠., ٥		- , -	395	0,0			-,0	400
	Ala	Phe	Ser	Glu		Ser	Arg	Leu	Arg		His	G1n	Arg	Thr	
, ,				405	,		- 0		410				3	415	
Thr	Clv	Clu	Lvc		Tyr	Lvc	Cvc	Acn		Cyc	Acn	Lvc	Δla		Sor

			420					425					430		
Ala	Lys	Ser	Gly	Leu	Arg	Ile	His	Gln	Arg	Thr	His	Thr	Gly	Glu	Lys
		435					440					445			
Pro	Phe	Glu	Cys	His	Glu	Cys	Gly	Lys	Ser	Phe	Asn	Tyr	Lys	Ser	Ile
	450					455					460				
Leu	He	Val	His	Gln	Arg	Thr	His	Thr	Gly	Glu	Lys	Pro	Phe	Glu	Cys
465					470					475					480
Asn	Glu	Cys	Gly	Lys	Ser	Phe	Ser	His	Met	Ser	Gly	Leu	Arg	Asn	His
				485					490					495	
Arg	Arg	Thr	His	Thr	Gly	Glu	Arg	Pro	Tyr	Lys	Cys	Asp	Glu	Cys	Gly
			500					505					510		
Lys	Ala	Phe	Lys	Leu	Lys	Ser	Gly	Leu	Arg	Lys	His	His	Arg	Thr	His
		515				٠	520					525			
Thr	G1y	Glu	Lys	Pro	Tyr	Lys	Cys	Asn	Gln	Cys	Gly	Lys	Ala	Phe	Gly
	530					535					540				
Gln	Lys	Ser	Gln	Leu	Arg	Gly	His	His	Arg	Ile	His	Thr	Gly	Glu	Lys
545					550					555					560
Pro	Tyr	Lys	Cys	Asn	His	Cys	Gly	Glu	Ala	Phe	Ser	Gln	Lys	Ser	Asn
				565					570					575	
Leu	Arg	Val	His	His	Arg	Thr	His	Thr	Gly	Glu	Lys	Pro	Tyr	Gln	Cys
			580					585					590		
Glu	Glu	Cys	Gly	Lys	Thr	Phe	Arg	Gln	Lys	Ser	Asn	Leu	Arg	Gly	His
		595					600					605			
Gln	Arg	Thr	His	Thr	Gly	Glu	Lys	Pro	Tyr	Glu	Cys	Asn	Glu	Cys	G1 y
	610					615					620				
Lys	Ala	Phe	Ser	Glu	Lys	Ser	Val	Leu	Arg	Lys	His	Gln	Arg	Thr	His
625					630					635					640
Thr	Gly	Glu	Lys	Pro	Tyr	Asn	Cys	Asn	Gln	Cys	Gly	Glu	Ala	Phe	Ser
				645					650					655	
Gln	Lys	Ser	Asn	Leu	Arg	Val	His	Gln	Arg	Thr	His	Thr	Gly	Glu	Lys
			660					665					670		
Pro	Tyr	Lys	Cys	Asp	Lys	Cys	Gly	Arg	Thr	Phe	Ser	Gln	Lys	Ser	Ser
		675					680					685			
Leu	Arg	Glu	His	Gln	Lys	Ala	His	Pro	Gly	Asp					
	690					695									

(

```
<211> 319
<212> PRT
<213> Homo sapiens
<400> 547
Met Glu Tyr Pro Ala Pro Ala Thr Val Gln Ala Ala Asp Gly Gly Ala
                  5
 1
                                      10
                                                          15
Ala Gly Pro Tyr Ser Ser Ser Glu Leu Leu Glu Gly Gln Glu Pro Asp
             20
                                 25
Gly Val Arg Phe Asp Arg Glu Arg Ala Arg Arg Leu Trp Glu Ala Val
                             40
                                                 45
Ser Gly Ala Gln Pro Val Gly Arg Glu Glu Ala Gln Phe Pro His
    50
                         55
Ser Arg Thr Val Ile Pro Ile Leu Val Leu Ser Glu Thr Tyr Ser Leu
                     70
                                          75 ·
Cys His Pro Val Glu His Met Ile Gln Lys Asn Gln Cys Leu Phe Thr
                 85
                                     90
                                                          95
Asn Thr Gln Cys Lys Val Cys Cys Ala Leu Leu Ile Ser Glu Ser Gln
            100
                                105
                                                     110
Lys Leu Ala His Tyr Gln Ser Lys Lys His Ala Asn Lys Val Lys Arg
                            120
                                                 125
Tyr Leu Ala 11e His Gly Met Glu Thr Leu Lys Gly Glu Thr Lys Lys
    130
                        135
Leu Asp Ser Asp Gln Lys Ser Ser Arg Ser Lys Asp Lys Asn Gln Cys
                    150
                                         155
Cys Pro Ile Cys Asn Met Thr Phe Ser Ser Pro Val Val Ala Gln Ser
                                     170
                                                         175
                165
His Tyr Leu Gly Lys Thr His Ala Lys Asn Leu Lys Leu Lys Gln Gln
            180
                                185
                                                     190
Ser Thr Lys Val Glu Ala Leu His Gln Asn Arg Glu Met Ile Asp Pro
                            200
Asp Lys Phe Cys Ser Leu Cys His Ala Thr Phe Asn Asp Pro Val Met
    210
                        215
                                             220
```

Ala Gln Gln His Tyr Val Gly Lys Lys His Arg Lys Gln Glu Thr Lys

<210> 4924

225 230 235 240 Leu Lys Leu Met Ala Arg Tyr Gly Arg Leu Ala Asp Pro Ala Val Thr 245 250 255 Asp Phe Pro Ala Gly Lys Gly Tyr Pro Cys Lys Thr Cys Lys Ile Val 260 265 270 Leu Asn Ser Ile Glu Gln Tyr Gln Ala His Val Ser Gly Phe Lys His 280 Lys Asn Gln Ser Pro Lys Thr Val Ala Ser Ser Leu Gly Gln Ile Pro 290 295 300 Met Gln Arg Gln Pro Ile Gln Lys Asp Ser Thr Thr Leu Glu Asp 305 310 315

<210> 4925

<211> 458

<212> PRT

<213> Homo sapiens

<400> 548

Met Glu Ser Asn Phe Asn Thr Glu Ser Ser Ser Thr Phe Thr Leu Gln 10 Ser Ser Ser Glu Thr Leu Phe Ser Ile Gln Leu Leu Asp Phe Lys Thr 20 25 Ser Leu Leu Glu Ala Leu Glu Glu Leu Arg Met Arg Arg Glu Ala Glu 35 40 45 lle His Tyr Glu Glu Gln lle Gly Lys lle lle Val Glu Thr Gln Glu 55 Leu Lys Trp Gln Lys Glu Thr Leu Gln Asn Gln Lys Glu Thr Leu Ala 65 70 75 80 Glu Gln His Lys Glu Ala Met Ala Val Phe Lys Lys Gln Leu Gln Met 85 90 Lys Met Cys Ala Leu Glu Glu Glu Lys Gly Lys Tyr Gln Leu Ala Thr

100 105 110

Glu 11e Lys Glu Lys Glu 11e Glu Gly Leu Lys Glu Thr Leu Lys Ala 115 120 125

Leu Gln Val Ser Lys Tyr Ser Leu Gln Lys Lys Val Ser Glu Met Glu

	130					135					140				
Gln	Lys	Val	Gln	Leu	His	Leu	Leu	Ala	Lys	Glu	Asp	Tyr	His	Lys	Gln
145					150					155					160
Leu	Ser	Glu	11e	Glu	Lys	Tyr	Tyr	Ala	Thr	He	Thr	Gly	Gln	Phe	Gly
				165					170					175	
Leu	Val	Lys	Glu	Asn	His	Glu	Lys	Leu	Glu	Gln	Asn	Val	Arg	Glu	Ala
			180					185					190		
He	Gln	Ser	Asn	Lys	Arg	Leu	Ser	Ala	Leu	Asn	Lys	Lys	Gln	Glu	Ala
		195					200					205			
Glu	lle	Cys	Ser	Leu	Lys	Lys	Glu	Leu	Lys	Lys	Ala	Ala	Ser	Asp	Leu
	210					215					220				
He	Lys	Ser	Lys	Val	Thr	Cys	Gln	Tyr	Lys	Met	Gly	Glu	Glu	Asn	He
225					230					235					240
Asn	Leu	Thr	He	Lys	Glu	Gln	Lys	Phe	Gln	Glu	Leu	Gln	Glu	Arg	Leu
				245					250					255	
Asn	Met	Glu	Leu	Glu	Leu	Asn	Glu	Lys	He	Asn	Glu	Glu	He	Thr	His
			260					265					270		
He	Gln	Glu	Glu	Lys	Gln	Asp	lle	Ile	Ile	Ser	Phe	Gln	His	Met	Gln
		275					280					285			
Gln		Leu	Arg	Gln	Gln	Ile	Gln	Ala	Asn	Thr	Glu	Met	Glu	Ala	Glu
	290					295					300				
Leu	Lys	Val	Leu	Lys	Glu	Asn	Asn	Gln	Thr	Leu	Glu	Arg	Asp	Asn	
305					310					315					320
Leu	Gln	Arg	Glu		Val	Lys	Glu	Asn		Glu	Lys	Phe	Leu		Leu
				325					330	_	_			335	
Gln	Asn	Glu		Glu	Lys	Ala	Leu		Thr	Trp	Lys	Arg		Ala	Glu
			340	0.1	7.1			345			0.1		350		
GIu	Leu		Gly	Glu	He	Asn		He	Lys	Asn	Glu		Ser	Ser	Leu
	0.1	355			0.1		360	61		æ.		365			
Lys		Ihr	HIS	He	Glu	Leu	GIn	Glu	HIS	lyr		Lys	Leu	Cys	Asn
C1	370	ть	DI	C I	C1	375	,	1	DI.	C1	380	17 1	D	C1	V 1
	Lys	Inr	Phe	Glu		Asp	Lys	Lys	Phe		Asn	Val	Pro	61u	
385	Λ ~ ~	Clu	A 65	C ~	390	Mo *	C.	TL	C1	395	C ~ ~	C1.	۸	T1	400
ASII	ASII	010	ASII		610	Met	ser	mr		Lys	ser	GIU	ASN		116
116	Gla	lve	Tun	405	The	Glu	Cla	Gla	410	Ara	Cl.	61	Acr	415 Mot	Cl.
110	OIL	1.V.S	1 V I	ASH	1111	11111	11111	11111	1 1 1-4	niy	11111		M > 11	WILL-F I	

Asn Phe Cys Ser Asp Thr Glu Tyr Arg Glu Lys Glu Glu Lys Lys Arg Arg Leu Ile Tyr Arg Gly Asn Asn Tyr Arg <210> 4926 <211> 814 <212> PRT <213> Homo sapiens <400> 549 Met Val Lys Gln Thr 11e Gln 11e Phe Ala Arg Val Lys Pro Pro Val Arg Lys His Gln Gln Gly Ile Tyr Ser Ile Asp Glu Asp Glu Lys Leu Ile Pro Ser Leu Glu Ile Ile Leu Pro Arg Asp Leu Ala Asp Gly Phe Val Asn Asn Lys Arg Glu Ser Tyr Lys Phe Lys Phe Gln Arg Ile Phe Asp Gln Asp Ala Asn Gln Glu Thr Val Phe Glu Asn Ile Ala Lys Pro Val Ala Gly Ser Val Leu Ala Gly Tyr Asn Gly Thr lle Phe Ala Tyr Gly Gln Thr Gly Ser Gly Lys Thr Phe Thr 11e Thr Gly Gly Ala Glu Arg Tyr Ser Asp Arg Gly Ile Ile Pro Arg Thr Leu Ser Tyr Ile Phe Glu Gln Leu Gln Lys Asp Ser Ser Lys Ile Tyr Thr Thr His Ile Ser Tyr Leu Glu lle Tyr Asn Glu Cys Gly Tyr Asp Leu Leu Asp Pro Arg His Glu Ala Ser Ser Leu Glu Asp Leu Pro Lys Val Thr 11e Leu Glu

Asp Pro Asp Gln Asn Ile His Leu Lys Asn Leu Thr Leu His Gln Ala

			180					185					190		
Thr	Thr	Glu	Glu	Glu	Ala	Leu	Asn	Leu	Leu	Phe	Leu	Gly	Asp	Thr	Asn
		195					200					205			
Arg	Met	He	Ala	Glu	Thr	Pro	Met	Asn	Gln	Ala	Ser	Thr	Arg	Ser	His
	210					215					220				
Cys	Île	Phe	Thr	He	His	Leu	Ser	Ser	Lys	Glu	Pro	Gly	Ser	Ala	Thr
225					230					235					240
Val	Arg	His	Ala	Lys	Leu	His	Leu	Val	Asp	Leu	Ala	Gly	Ser	Glu	Arg
				245					250					255	
Val	Ala	Lys	Thr	Gly	Val	G1y	Gly	His	Leu	Leu	Thr	Glu	Ala	Lys	Tyr
			260					265					270		
He	Asn	Leu	Ser	Leu	His	Tyr	Leu	Glu	Gln	Val	He	lle	Ala	Leu	Ser
		275					280					285			
Glu		His	Arg	Ser	His	He	Pro	Tyr	Arg	Asn	Ser	Met	Met	Thr	Ser
	290					295					300				
	Leu	Arg	Asp	Ser		Gly	Gly	Asn	Cys		Thr	Thr	Met	He	
305					310					315					320
Thr	Leu	Ser	Leu		Lys	Arg	Asn	Leu		Glu	Ser	He	Ser		Cys
	101	. 1	0.1	325	1			T.	330		0.1			335	
Arg	Phe	Ala		Arg	Val	Ala	Leu		Lys	Asn	GIu	Ala		Leu	Asn
61	61	7.1	340	D		,		345				0.1	350	6.1	
Glu	Glu		Asn	Pro	Arg	Leu	Val	lle	Lys	Arg	Leu		Lys	Glu	11e
C1	C1	355	1	Α	C1	1	360	Mad	V-1	Tl	C1	365	C1	A	Th
GIN		Leu	Lys	ASP	Glu		Ala	we t	vai	1 1113-		Glu	GIN	Arg	ınr
Clu	370	Lou	Thr	61	A 1 o	375	Leu	Lou	Cln	Lou	380	Lvc	Lou	110	The
385	пта	Leu	1111	Giu	390		Leu	Leu	OIII	395		Lys	Leu	116	400
	Phe	Leu	Glu	Asn	-		Ser	Asn	Ser			Glu	Val	Glv	
50,	7 110	Lea	oru	405	OIII	пор	501	пор	410	, 11 E	Leu	oru	741	415	711 G
Asp	Met	Arg	Lvs		His	His	Cys	Phe		His	l.eu	Lvs	Lvs		Leu
,		0	420					425				-,-	430		
Asn	Asp	Lys		lle	Leu	Glu	Asn		Thr	Val	Ser	Ser	Glu	Ser	Lys
		435	•				440					445			
Asp	Gln	Asp	Cys	G1n	Glu	Pro	Leu	Lys	Glu	Glu	Glu		Arg	Lys	Leu
	450					455					460				
Arg	Asp	He	Leu	Lvs	Gln	Arg	Asp	Asn	Glu	He	Asn	lle	Leu	Val	Asn

465					470					475					480
Met	Leu	Lys	Lys	Glu	Lys	Lys	Lys	Ala	Gln	Glu	Ala	Leu	His	Leu	Ala
				485					490					495	
Gly	Met	Asp	Arg	Arg	Glu	Phe	Arg	Gln	Ser	Gln	Ser	Pro	Pro	Phe	Arg
			500					505					510		
Leu	Gly	Asn	Pro	Glu	Glu	Gly	Gln	Arg	Met	Arg	Leu	Ser	Ser	Ala	Pro
		515					520					525			
Ser	Gln	Ala	Gln	Asp	Phe	Ser	Ile	Leu	Gly	Lys	Arg	Ser	Ser	Leu	Leu
	530					535					540				
His	Lys	Lys	Ile	G1 y	Met	Lys	Glu	Glu	Met	Ser	Leu	Gly	Cys	Gln	Glu
545					550					555					560
Ala	Phe	Glu	He	Phe	Lys	Arg	Asp	His	Ala	Asp	Ser	Va]	Thr	He	Asp
				565					570					575	
Asp	Asn	Lys	Gln	He	Leu	Lys	Gln	Arg	Phe	Ser	Glu	Ala	Lys	Ala	Leu
			580					585					590		
Gly	Glu	Ser	lle	Asn	Glu	Ala	Arg	Ser	Lys	Ile	Gly	His	Leu	Lys	Glu
		595					600					605			
Glu	lle	Thr	Gln	Arg	His	lle	Gln	Gln	Val	Ala	Leu	Gly	Ile	Ser	Glu
	610					615					620				
Asn	Met	Ala	Val	Pro	Leu	Met	Pro	Asp	Gln	Gln	Glu	Glu	Lys	Leu	Arg
625					630					635					640
Ser	Gln	Leu	Glu	Glu	Glu	Lys	Arg	Arg	Tyr	Lys	Thr	Met	Phe	Thr	Arg
				645					650					655	
Leu	Lys	Ala	Leu	Lys	Val	Glu	lle	Glu	His	Leu	Gln	Leu	Leu	Met	Asp
			660					665					670		
Lys	Ala	Lys	Val	Lys	Leu	Gln	Lys	Glu	Phe	Glu	Val	Trp	Trp	Ala	Glu
		675					680					685			
Glu	Ala	Thr	Asn	Leu	Gln	Val	Asn	Ser	Pro	Ala	Val	Asn	Ser	Leu	Asp
	690					695					700				
His	Thr	Lys	Pro	Phe	Leu	Gln	Thr	Ser	Asp	Phe	Gln	His	Glu	Arg	Ser
705					710					715					720
Gln	Leu	Leu	Ser	Asn	Lys	Ser	Ser	Gly	Gly	Trp	Glu	Val	Gln	Asp	Gln
				725					730					735	
Gly	Thr	Gly	Arg	Phe	Asp	Val	Cys	Asp	Va1	Asn	Ala	Arg	Lys	He	Leu
			740					745					750		
Pro	Ser	Pro	Cvs	Pro	Ser	Pro	Hie	Ser	G1n	Lvc	Gln	Ser	Ser	Thr	Ser

Thr Pro Leu Gly Asp Ser Ile Pro Lys Arg Pro Val Ser Ser Ile Pro Leu Thr Gly Asp Ser Gln Thr Asp Ser Asp lle lle Ala Phe Ile Lys Ala Arg Gln Ser lle Leu Gln Lys Gln Cys Leu Gly Ser Asn <210> 4927

<211> 503 <212> PRT <213> Homo sapiens

<400> 550

Met Gly Ala Leu Thr Phe Arg Asp Val Ala Ile Glu Phe Ser Leu Glu Glu Trp Gln Cys Leu Asp Thr Glu Gln Gln Asn Leu Tyr Arg Asn Val Met Leu Asp Asn Tyr Arg Asn Leu Val Phe Leu Gly Ile Ala Val Ser

Lys Pro Asp Leu Ile Thr Cys Leu Glu Glu Glu Lys Glu Pro Trp Asn

Leu Lys Thr His Asp Met Val Ala Lys Pro Pro Val Ile Cys Ser His

Ile Ala Gln Asp Leu Trp Pro Glu Gln Gly Ile Lys Asp Tyr Phe Gln

Glu Val 11e Leu Arg Gln Tyr Lys Lys Cys Arg His Glu Asn Leu Leu

Leu Arg Lys Gly Cys Lys Asn Val Asp Glu Phe Lys Met His Lys Lys

Gly Tyr Asn Arg His Asn Gln Cys Leu Thr Thr Ser His Ser Lys 11e

Phe Gln Cys Asp Lys Tyr Val Lys Val Phe His Lys Phe Ser Asn Ser

Asn Arg His Lys lle Arg His Thr Ser Lys Lys Pro Phe Lys Cys Lys

			165					170					175	
Cys	Gly	Lys	Leu	Phe	Cys	He	Leu	Ser	His	Leu	Ala	Gln	His	Lys
		180					185					190		
Пе	llis	Thr	Gly	Glu	Lys	Ser	Tyr	Lys	Cys	Glu	Glu	Tyr	Gly	Lys
	195					200					205			
Phe	Asn	Glu	Ser	Ser	Asn	Cys	Thr	Thr	His	Lys	Arg	11e	Thr	Glu
210					215					220				
Lys	Pro	Tyr	Lys	Cys	Lys	Glu	Cys	Gly	Lys	Ala	Phe	Asn	Trp	Phe
				230					235					240
His	Phe	Thr	Thr	His	Lys	Arg	Ile	His	Thr	Gly	Glu	Lys	Pro	Tyr
			245					250					255	
Cys	Glu	Lys	Cys	Gly	Lys	Phe	Phe	Așn	Gln	Ser	Thr	Asn	Leu	Thr
		260					265					270		
His	Lys	Arg	He	llis	Thr	Gly	Glu	Lys	Pro	Tyr	Lys	Cys	Glu	Glu
	275					280					285			
Gly	Lys	Ala	Phe	Asn	Gln	Ser	Ser	Asn	Leu	Thr	Glu	His	Lys	Lys
290					295					300				
His	Thr	Lys	Glu	Gln	Pro	Tyr	Lys	Cys	Glu	Lys	Cys	Gly	Lys	Ala
				310					315					320
Lys	Trp	Ser	Ser	Thr	Leu	Thr	Lys	His	Lys	Arg	He	His	Asn	Gly
			325					330					335	
Lys	Pro	Tyr	Lys	Cys	Glu	Glu	Cys	G1 y	Lys	Ala	Phe	Asn	Arg	Ser
		340					345					350		
Thr	Leu	Asn	Arg	His	Lys	Лe	Thr	His	Thr	Gly	Gly	Lys	Pro	Tyr
	355					360								
Tyr	Lys	Glu	Cys	Gły	Lys	Λla	Phe	Asn	Gln		Ser	Thr	Leu	Thr
370														
His	Lys	He	He		Thr	Val	GJu	Lys		Tyr	Lys	Cys	Glu	
			F 31							m	æ.			400
Gly	Lys	Ala		Ser	Arg	He	Ser		Leu	lhr	lhr	HIS		Arg
	mı.	0.1			ъ	T			C.1	6.1		61		. 1
HIS	Ihr		Glu	Lys	Pro	lyr		Cys	61u	61u	Cys		Arg	ATa
٨	C1.		C	TI	1 .	ть.		11:	1	Δ	11.		ть.	C1
Asn		ser	ser	ınr	ren		ını	nis	Lys	Arg		mis	ınr	оту
Luc		Tun	61	Cva	Clos		Cvc	Clv	Lve	A10		Acr	Arc	Sor
	Phe 210 Lys His Cys His Lys Lys Thr Tyr 370 His Gly His	Ile His 195 Phe Asn 210 Lys Pro His Phe Cys Glu His Lys 275 Gly Lys 290 His Thr Lys Trp Lys Pro Thr Leu 355 Tyr Lys 370 His Lys Gly Lys 370 His Lys Gly Lys	180 11e 11is Thr 195 Phe Asn Glu 210 Lys Pro Tyr 260 His Lys Arg 275 Gly Lys Ala 290 His Thr Lys Lys Pro Tyr 340 Thr Leu Asn 355 Tyr Lys Glu 370 His Lys Jle Gly Lys Ala Ala His Thr Gly 420 Asn Gln Ser 435	Cys Gly Lys Leu 11e His Thr Gly 195 - - Phe Asn Glu Ser 210 - Lys Lys Lys Pro Thr Thr 245 Cys Glu Lys Cys 260 -	Cys Gly Lys Leu Phe 11e His Thr Gly Glu 195 - Ser Ser 210 - Lys Cys Lys Pro Tyr Lys Cys Lys Phe Thr His 245 Cys Glu Lys Cys Gly Lys Arg Ile His 275 - 310 Lys Ala Phe Asn 290 - 310 His Thr Lys Glu Gln 11e His 310 Lys Trp Ser Thr 325 - Thr Lys Fro Tyr Lys Cys Tyr Lys Glu Cys Gly 370 - - 390 Gly Lys Ala Phe Ser His Lys Ala Phe Ser His	Cys Lys Leu Phe Cys 11e His Thr Gly Glu Lys 195	Cys Lys Leu Phe Cys 11e 11e 13o	Cys Gly Lys Leu Phe Cys 11e Leu 11e 11is Thr Gly Glu Lys Ser Tyr 11e 11is Thr Gly Glu Lys Ger Tyr 11e Asn Glu Ser Ser Asn Cys Thr 210 Tyr Lys Cys Lys Glu Cys Lys Pro Tyr Lys Cys Lys Arg Ile Lys Pro Tyr Lys Gly Lys Arg Ile Cys Glu Lys Cys Gly Lys Arg Ile His Lys Arg Ile Ilis Thr Gly Gly Gly Gly Gly Gly Gly Gly Gly Lys Ser Ser Ser Ser Gly Lys Lys Lys Arg Ile Lys Arg Ile Ile Ile Ile Ile Ile Ile <t< td=""><td>Cys Gly Lys Leu Phe Cys 11e Leu Ser 11e His Thr Gly Glu Lys Ger Tyr Lys 11e His Thr Gly Gu Lys Gy Tyr Lys 195 Tr Tr Ger Asn Cys Thr Thr 210 Tr Tyr Lys Cys Lys Gu Thr Thr 11e Pro Tyr Lys Cys Lys Gly Cys Gly Gly<td>Cys Gly Lys Leu Phe Cys 11e Leu Ser His 11e His Thr Gly Glu Lys Ser Tyr Lys Cys 11e His Thr Gly Glu Lys Ser Tyr Lys Cys 11e Asn Glu Ser Ser Asn Cys Thr His 210 Tr Lys Cys Lys Glu Cys Gly Lys Gl</td><td>Cys Gly Lys Leu Phe Cys 11e Leu Ser His Leu 11e 11s Thr Gly Gly Lys Ser Tyr Lys Cys Glu 11e Asn Gly Ser Asn Lys Thr Hys Cys Gly Lys Thr Hys Lys Lys Thr Hys Lys Asn Lys Lys Ana 230 Lys Lys Lys Ana 230 Lys Ana 235 Lys Ana 235 Lys Ana Ana 235 Lys Ana A</td><td>Cys Gly Lys Leu Phe Cys 11e Leu Ser His Leu Ala 11e 11s Thr Gly Glu Lys Ser Tyr Lys Cys Glu Glu Glu Glu Jus Jus Lys Lys Lys Tyr Lys Lys Arg Jus Lys L</td><td>Cys Gly Lys Leu Phe Cys 11e Leu His Leu Jab 190 11e His Thr Glu Glu Lys Ser Tyr Lys Glu Glu Tyr 11e His Thr Glu Glu Lys Asn Glu Lys Glu Lys Lys Lys Jab Jab</td><td>Cys Gys Leu Phes Cys 11e Leu Ser His Leu Ala His 11e His Thr Gly Gly Lys Ser Tyr Lys Cys Glu Glu Tyr Gly 11e His Thr Gly Gly Lys Gly Lys Gly Gly Lys Arg Gly Gly Lys Arg His Tyr Lys Arg His His Lys Arg His His</td></td></t<>	Cys Gly Lys Leu Phe Cys 11e Leu Ser 11e His Thr Gly Glu Lys Ger Tyr Lys 11e His Thr Gly Gu Lys Gy Tyr Lys 195 Tr Tr Ger Asn Cys Thr Thr 210 Tr Tyr Lys Cys Lys Gu Thr Thr 11e Pro Tyr Lys Cys Lys Gly Cys Gly Gly <td>Cys Gly Lys Leu Phe Cys 11e Leu Ser His 11e His Thr Gly Glu Lys Ser Tyr Lys Cys 11e His Thr Gly Glu Lys Ser Tyr Lys Cys 11e Asn Glu Ser Ser Asn Cys Thr His 210 Tr Lys Cys Lys Glu Cys Gly Lys Gl</td> <td>Cys Gly Lys Leu Phe Cys 11e Leu Ser His Leu 11e 11s Thr Gly Gly Lys Ser Tyr Lys Cys Glu 11e Asn Gly Ser Asn Lys Thr Hys Cys Gly Lys Thr Hys Lys Lys Thr Hys Lys Asn Lys Lys Ana 230 Lys Lys Lys Ana 230 Lys Ana 235 Lys Ana 235 Lys Ana Ana 235 Lys Ana A</td> <td>Cys Gly Lys Leu Phe Cys 11e Leu Ser His Leu Ala 11e 11s Thr Gly Glu Lys Ser Tyr Lys Cys Glu Glu Glu Glu Jus Jus Lys Lys Lys Tyr Lys Lys Arg Jus Lys L</td> <td>Cys Gly Lys Leu Phe Cys 11e Leu His Leu Jab 190 11e His Thr Glu Glu Lys Ser Tyr Lys Glu Glu Tyr 11e His Thr Glu Glu Lys Asn Glu Lys Glu Lys Lys Lys Jab Jab</td> <td>Cys Gys Leu Phes Cys 11e Leu Ser His Leu Ala His 11e His Thr Gly Gly Lys Ser Tyr Lys Cys Glu Glu Tyr Gly 11e His Thr Gly Gly Lys Gly Lys Gly Gly Lys Arg Gly Gly Lys Arg His Tyr Lys Arg His His Lys Arg His His</td>	Cys Gly Lys Leu Phe Cys 11e Leu Ser His 11e His Thr Gly Glu Lys Ser Tyr Lys Cys 11e His Thr Gly Glu Lys Ser Tyr Lys Cys 11e Asn Glu Ser Ser Asn Cys Thr His 210 Tr Lys Cys Lys Glu Cys Gly Lys Gl	Cys Gly Lys Leu Phe Cys 11e Leu Ser His Leu 11e 11s Thr Gly Gly Lys Ser Tyr Lys Cys Glu 11e Asn Gly Ser Asn Lys Thr Hys Cys Gly Lys Thr Hys Lys Lys Thr Hys Lys Asn Lys Lys Ana 230 Lys Lys Lys Ana 230 Lys Ana 235 Lys Ana 235 Lys Ana Ana 235 Lys Ana A	Cys Gly Lys Leu Phe Cys 11e Leu Ser His Leu Ala 11e 11s Thr Gly Glu Lys Ser Tyr Lys Cys Glu Glu Glu Glu Jus Jus Lys Lys Lys Tyr Lys Lys Arg Jus Lys L	Cys Gly Lys Leu Phe Cys 11e Leu His Leu Jab 190 11e His Thr Glu Glu Lys Ser Tyr Lys Glu Glu Tyr 11e His Thr Glu Glu Lys Asn Glu Lys Glu Lys Lys Lys Jab Jab	Cys Gys Leu Phes Cys 11e Leu Ser His Leu Ala His 11e His Thr Gly Gly Lys Ser Tyr Lys Cys Glu Glu Tyr Gly 11e His Thr Gly Gly Lys Gly Lys Gly Gly Lys Arg Gly Gly Lys Arg His Tyr Lys Arg His His Lys Arg His His

Ser Thr Leu Thr Thr His Lys Ile Ile His Ser Gly Glu Lys lle Tyr Lys Cys Lys Glu Cys Gly Lys Ala Phe Arg Arg Phe Ser His Leu Thr Arg His Lys Thr 11e His Thr <210> 4928 <211> 388 <212> PRT <213> Homo sapiens <400> 551 Met Leu Leu Lys Glu Lys Glu Asp Ser Leu Met Thr Cys Gln Gln Ile Tyr Lys Ala Leu Gln Glu Glu Leu Thr Val Lys Glu Lys Gln Glu Glu Asp lle Lys Arg Arg lle Asn Leu Ala Glu Asn Glu Leu Glu lle Thr Lys Thr Leu Leu Asn Gln Thr Arg Glu Glu Val Leu Thr Leu Lys Asn Glu Arg Glu Leu Met Leu Ile Ser His Gln Lys Ser Ile Glu Gln Leu Gln Glu Thr Leu Arg Gln Lys Leu Leu Ser Asp Asp Asn Trp Lys Glu Lys 11e Glu Ala Glu Leu Ala Lys Glu Arg Ala Gln His Leu Val Glu Phe Glu Glu Gln Ala Leu Leu Phe Lys Glu Glu Thr Lys Leu Gln Leu Asp Ile Glu Lys Glu Lys His Gln Asp Val Ile Gln Lys Tyr Lys Lys Glu Gln Glu Glu Leu Gln Met Lys lle Ser Asp Leu lle Thr Gly Ala

Thr Arg Asp Leu Arg Gln Glu Val Thr Thr Leu Lys Glu Lys Leu His

				165					170					175	
Lys	Ser	His	Thr	Arg	Tyr	Thr	Glu	Glu	Ser	Asn	Ser	Lys	Glu	Lys	Glu
			180					185					190		
lle	Glu	Asn	Leu	Lys	Asn	Leu	Val	Ala	G] u	Phe	Glu	Ser	Arg	Leu	Lys
		195					200					205			
Lys	Glu	He	Лѕр	Ser	Asn	Asp	Ser	Val	Ser	Glu	Asn	Leu	Arg	Lys	Glu
	210					215					220				
Met	Glu	Gln	Lys	Ser	Asp	Glu	Leu	Lys	Arg	Val	Met	Leu	Ala	Gln	Thr
225					230					235					240
Gln	Leu	He	Glu	Gln	Phe	Asn	Gln	Ser	Gln	Glu	Glu	Asn	Thr	Phe	Leu
				245					250					255	
Gln	Glu	Thr	Val	Arg	Arg	Glu	Cys	Glu	Glu	Arg	Phe	Ğlu	Leu	Thr	Glu
			260					265					270		
Ala	Leu	Ser	Gln	Ala	Arg	Glu	Gln	Leu	Leu	Glu	Leu	Ser	Lys	Leu	Arg
		275					280					285			
Gly	Ser	Leu	Pro	Phe	Ser	Pro	Cys	Ser	Leu	Ser	Lys	Gly	Ser	Leu	Thr
	290					295					300				
Ser	Pro	Ala	Ala	Ala	Val	Ser	Asn	His	Gly	Glu	Arg	Ser	Leu	Ala	Arg
305					310					315					320
Leu	Asn	Ser	Glu	Lys	Gly	Ile	Gln	lle	Pro	Asn	Leu	Arg	Gly	Val	Ser
				325		•			330					335	
Lys	Pro	Thr	Thr	Phe	Pro	Thr	Ser	Asp	Lys	Pro	Lys	Arg	Va]	Arg	Ser
			340					345					350		
Gly	Val	Pro	He	Leu	Pro	Gln	Pro	His	Pro	Pro	Arg	G1 y	Gly	Ala	Ser
		355					360					365			
Ser	Ala	Asn	Glu	Thr	Arg	Gln	Arg	Leu	Ala	Ala	lle	Leu	Arg	Arg	Arg
	370					375					380				
Arg	Ser	Gln	Gln												
385															

<211> 165

<212> PRT

<213> Homo sapiens

<400> 552 Met Asp Gly Pro Ala Thr Pro Val Ser Thr Asp Ser Asn Pro Pro Thr Gln Gln Glu Asp Arg Ser Ala Cys Lys Cys Thr His Leu Glu Lys Arg Leu Phe Pro Leu Leu Leu Val Ala Gln Leu Leu Leu Ser Pro Pro Gly Ala Ala Ala Val Lys Cys Gln Leu Asp Pro Ala Lys Trp Gln Asp Pro Gln His Ser Ser Ile Cys Ser Val Leu His Leu Arg His Trp Lys Gly Cys Glu Pro Asp Ile Gly Ser Gln Ser Thr Cys Phe Pro Glu Pro Glu Ser Cys Leu Pro Val Ala Ala Asp Thr Asp Ser Asn Val Thr Pro Ala Thr Gln Gln Gln Arg Cys Cys Thr Leu Ala Cys Asn Leu Gly Thr Gly Pro Leu His Leu Leu Leu Ser Leu Leu Met Gln Leu Gly Ala Arg Ala Cys Ala Thr Gly Ser Asp Leu Thr Ser Thr Ser Ser Arg Ala Thr Val Asn Leu His Val Pro <210> 4930 <211> 832 <212> PRT <213> Homo sapiens <400> 553 Met Ala Gly Leu Arg Tyr Ser Val Lys Val Tyr Val Leu Asn Glu Asp Glu Glu Trp Asn Asn Leu Gly Thr Gly Gln Val Ser Ser Thr Tyr Asp

Glu	Gln	Phe	Gln	Gly	Met	Ser	Leu	Leu	Val	Arg	Ser	Asp	Ser	Asp	Gly
		35					40					45			
Ser	Val	He	Leu	Arg	Ser	Gln	Ile	Pro	Pro	Asp	Arg	Pro	Tyr	Gly	Lys
	50					55					60				
Tyr	Gln	Glu	Thr	Leu	Ile	Val	Trp	Tyr	Glu	Ala	Glu	Asn	Gln	Gly	Leu
65					70					75					80
Val	Leu	Lys	Phe	Gln	Asp	Pro	Ala	Gly	Cys	Gln	Asp	lle	Trp	Lys	Glu
				85					90					95	
Ile	Cys	Gln	Ala	Gln	Gly	Lys	Asp	Pro	Ser	Ile	Gln	Thr	Thr	Val	Asn
			100					105					110		
Ile	Ser	Asp	Glu	Pro	Glu	Glu	Asp	Phe	Asn	Glu	Met	Ser	Val	He	Ser
		115					120					125			
Asn	Met	Val	Val	Leu	Pro	Asp	Cys	Glu	Leu	Asn	Thr	Leu	Asp	G]n	He
	130					135					140				
Ala	Asp	He	Val	Thr	Ser	Val	Phe	Ser	Ser	Pro	Val	Thr	Asp	Arg	Glu
145					150					155					160
Arg	Leu	Ala	Glu		Leu	Lys	Asn	Glu	Ala	Tyr	Ile	Pro	Lys	Leu	Leu
				165					170					175	
Gln	Leu	Phe		Thr	Cys	Glu	Asn		Glu	Asn	Thr	Glu	G1 y	Leu	His
		_	180			_		185					190		
His	Leu		Glu	He	He	Lys	Gly	He	Leu	Phe	Leu		Glu	Ala	Cys
	DI	195			D1	0	200	0.1				205			
Leu		Glu	He	Met	Phe		Asp	Glu	Cys	He		Asp	Val	Val	Gly
0	210	C1	т		Б	215			0.1	D	220				
	Leu	Glu	lyr	Asp		Ala	Leu	Asp	GIn		Lys	Arg	His	Arg	
225 DI		TI			230		DI		C1	235	7.7	Б	7.1	T)	240
rne	Leu	Inr	ASN		мта	Lys	Phe	Lys		val	116	Pro	11e		Asn
Cor	C1	Lau	A 200	245	Luc	I l a	Ui a	Cl.	250	Т	Λ	1	C1	255	т1.
ser	GJU	Leu		GIN	Lys	11e	His		ınr	lyr	Arg	Leu		lyr	116
Tun	Aan	110	260 Law	Lau	Dno	Vol.	Drag	265	Tla	Dha	C1	A ===	270	Dlag	1
I y I	nsp	275	Leu	Leu	110	vai	Pro	ser	116	rne	G1u		ASII	rne	Leu
Sor	Thr		The	Thr	Dho	Ha	280 Pho	Son	Acn	Lyo	Λlο	285	Ha	Vol.	Com
OCI	290	ı,cu	1 111	1111	i ne	295	Phe	261	Non	rys	300	Olu	116	101	Sel.
Met		Gla	lve	Ace	Hic		Phe	Leu	Tyr	Glo		Pho	Δla	Gla	Leu
305	.,, u	0111	1- y S	nap	310	LyS	HE	ren	1 y 1	315	· a I	ine	1110	0111	rea

Lys	Asp	Glu	Thr	Thr 325	His	Asp	Asp	Arg	Arg 330	Cys	Glu	Leu	Leu	Phe 335	Phe
Phe	Lys	Glu	Leu	Cys	Ser	Phe	Ser	Gln	Ala	Leu	Gln	Pro	Gln	Ser	Lys
			340					345					350		
Asp	Ala	Leu	Phe	Glu	Thr	Leu	Пе	Gln	Leu	Gl y	Val	Leu	Pro	Ala	Leu
		355					360					365			
Lys	Ile	Val	Met	Пе	Arg	Asp	Asp	Leu	Gln	Val	Arg	Ser	Ala	Ala	Ala
	370					375					380				
Val	Ile	Cys	Ala	Tyr	Leu	Val	Glu	Tyr	Ser	Pro	Ser	Arg	Ile	Arg	Glu
385					390					395					400
Phe	Ile	Ile	Ser	Glu	Ala	His	Val	Cys	Lys	Asp	Ile	Tyr	Leu	Phe	He
				405					410					415	
Asn	Val	11e	lle	Lys	Gln	Met	11e	Cys	Asp	Thr	Asp	Pro	Glu	Leu	Gly
			420					425					430		
Gly	Ala	Val	His	Leu	Met	Val	Val	Leu	His	Thr	Leu	Leu	Asp	Pro	Arg
		435					440					445			
Asn	Met	Leu	Thr	Thr	Pro	Glu	Lys	Ser	Glu	Arg	Ser	Glu	Phe	Leu	His
	450					455					460				
Phe	Phe	Tyr	Lys	His	Cys	Met	His	Lys	Phe	Thr	Ala	Pro	Leu	Leu	Ala
465					470					475					480
Ala	Thr	Ser	Glu	His	Asn	Cys	Glu	Glu	Asp	Asp	He	Ala	G1 y	Tyr	Asp
				485					490					495	
Lys	Ser	Lys	Asn	Cys	Pro	Asn	Asp	Asn	Gln	Thr	Ala	Gln	Leu	Leu	Ala
			500					505					510		
Leu	Ile	Leu	Glu	Leu	Leu	Thr	Phe	Cys	He	Gln	His	His	Thr	Phe	Tyr
		515					520					525			
He	Arg	Ser	Tyr	He	Leu	Asn	Lys	Asp	Leu	Leu	Arg	Lys	Ala	Leu	He
	530					535					540				
Leu	Met	Asn	Ser	Lys	His	Thr	His	Leu	He	Leu	Cys	Val	Leu	Arg	
545					550					555					560
Met	Arg	Arg	Met		Cys	Leu	Asn	Asp		Ala	Tyr	Asn	Asn	Tyr	He
			•	565					570					575	
He	Lys	G] y	Asn	Leu	Phe	Glu	Pro		Val	Asn	Ala	Leu	Leu	Asp	Asn
			580					585					590		
G1y	Thr		Tyr	Asn	Met	Leu	Asn	Ser	Ala	He	Leu		Leu	Phe	Glu
		595					600					605			

```
Tyr Ile Arg Val Glu Asn Ile Lys Pro Leu Val Ser His Ile Val Glu
                        615
                                            620
Lys Phe Tyr Asn Thr Leu Glu Ser Ile Glu Tyr Val Gln Thr Phe Lys
625
                    630
                                        635
                                                             640
Gly Leu Lys Ile Lys Tyr Glu Lys Glu Arg Asp Arg Gln Ser Gln Ile
                645
                                    650
Gln Lys Asn Leu His Ser Val Leu Gln Asn Ile Val Val Phe Arg Gly
                                665
Thr Ile Glu Glu Ile Gly Leu Glu Glu Glu Ile Cys Phe Met Glu Asp
                            680
Ala Gly Glu Ala Val Met Pro Pro Leu Glu Asp Asp Asp Glu Phe Met
                        695
                                            700
Glu Thr Lys Arg Thr Gln Glu Gly Glu Ala Val Met Pro Pro Leu Glu
705
                    710
                                        715
                                                             720
Asp Asp Asp Lys Phe Thr Glu Thr Lys Arg Thr His Gln Glu Gly Glu
                725
                                    730
Ala Val Met Pro Pro Leu Glu Asp Asp Asp Glu Phe Met Glu Thr Lys
                                745
Arg Asn Gln Glu His Glu Gly Lys Val Asp Ser Pro Lys Arg Thr Ser
        755
                            760
Ser Gly Asp Phe Lys Phe Ser Ser Ser Tyr Ser Ala Cys Ala Ala Ile
                        775
                                            780
Gly Thr Gly Ser Pro Ser Gly Ser Ser Val Val Arg Leu Val Asp His
                    790
                                        795
785
                                                             800
Pro Asp Asp Glu Glu Glu Lys Glu Glu Asp Glu Glu Glu Lys Glu Glu
                805
                                    810
Asp Lys Glu Asp Glu Thr Ser Pro Lys Lys Lys Pro His Leu Ser Ser
            820
                                825
                                                     830
```

<211> 250

<212> PRT

<213> Homo sapiens

<400> 554

Met	Phe	His	Gly	Thr	Val	Thr	Glu	Glu	Leu	Thr	Ser	His	Glu	Glu	Trp
1				5					10					15	
Ser	His	Tyr	Asn	Glu	Asn	He	Arg	Glu	Gly	Gln	Lys	Asp	Phe	Val	Phe
			20					25					30		
Val	Lys	Phe	Asn	Gly	Leu	His	Leu	Lys	Ser	Met	Glu	Asn	Leu	Gln	Ser
		35					40					45	•		
Cys	He	Ser	Leu	Arg	Val	Cys	He	Phe	Ser	Asn	Asn	Phe	lle	Thr	Asp
	50					55					60				
Ile	His	Pro	Leu	Gln	Ser	Cys	Ile	Lys	Leu	lle	Lys	Leu	Asp	Leu	His
65					70					75					80
Gly	Asn	Gln	He	Lys	Ser	Leu	Pro	Asn	Thr	Lys	Phe	Trp	Asn	Gly	Leu
				85					90					95	
Lys	Asn	Leu	Lys	Leu	Leu	Tyr	Leu	His	Λsp	Asn	Gly	Phe	Ala	Lys	Leu
			100					105					110		
Lys	Asn	lle	Cys	Val	Leu	Ser	Ala	Cys	Pro	Thr	Leu	He	Ala	Leu	Thr
		115					120					125			
Met	Phe	Asp	Cys	Pro	Val	Ser	Leu	Lys	Lys	Gly	Tyr	Arg	His	Val	Leu
	130					135					140				
Val	Asn	Ser	lle	Trp	Pro	Leu	Lys	Ala	Leu	Asp	His	His	Val	Ile	Ser
145					150					155					160
Asp	Glu	Glu	He	Ile	Gln	Asn	Trp	His	Leu	Pro	Glu	Arg	Phe	Lys	Ala
				165					170					175	
Cys	Asn	His	Arg	Leu	Phe	Phe	Asn	Phe	Cys	Pro	Ala	Leu	Arg	Lys	Gly
			180					185					190		
Thr	Thr	Tyr	G1u	Glu	Glu	lle	Asn	Asn	He	Lys	His	He	Thr	Ser	Lys
		195					200					205			
He	Asn	Λla	He	Leu	Ala	His	Asn	Ser	Pro	Val		He	Val	Gln	Arg
	210					215					220				
Trp	He	Arg	Gly	Phe	Leu	Val	Arg	Lys	Asn	Leu	Ser	Pro	Val	Phe	Phe
225					230					235					240
His	Lys	Lys	Lys	Thr	Ala	Gly	Lys	Asn	Tyr						
				245					250						

<211> 788

```
<213> Homo sapiens
<400> 555
Met Ala Cys Leu Thr His Arg Asn Glu Thr Asp Ala Arg Met Glu Phe
Tyr Ser Leu Phe His Lys Gly Asn Lys Ala Gly Val Gln Trp His Asp
                                 25
Leu Gly Ser Leu Gln Pro Leu Pro Pro Arg Phe Lys Arg Phe Ser Cys
                             40
                                                  45
Leu Ser Leu Gln Ser Ser Trp Asp Tyr Ser Leu Ser Lys Phe Asp Glu
                         55
                                              60
Arg Cys Cys Phe Leu Tyr Val His Asp Asn Ser Asp Asp Phe Gln 11e
65
                     70
                                          75
                                                              80
Tyr Phe Ser Thr Glu Glu Gln Cys Ser Arg Phe Phe Ser Leu Val Lys
                                     90
                 85
Glu Met Ile Thr Asn Thr Ala Gly Ser Thr Val Glu Leu Glu Gly Glu
                                105
Thr Asp Gly Asp Thr Leu Glu Tyr Glu Tyr Asp His Asp Ala Asn Gly
        115
                                                 125
Glu Arg Val Val Leu Gly Lys Gly Thr Tyr Gly Ile Val Tyr Ala Gly
                        135
Arg Asp Leu Ser Asn Gln Val Arg lle Ala lle Lys Glu lle Pro Glu
                    150
                                         155
                                                             160
145
Arg Asp Ser Arg Tyr Ser Gln Pro Leu His Glu Glu lle Ala Leu His
                165
                                     170
Lys Tyr Leu Lys His Arg Asn Ile Val Gln Tyr Leu Gly Ser Val Ser
                                185
Glu Asn Gly Tyr lle Lys Ile Phe Met Glu Gln Val Pro Gly Gly Ser
        195
                            200
                                                 205
Leu Ser Ala Leu Leu Arg Ser Lys Trp Gly Pro Met Lys Glu Pro Thr
                        215
                                             220
lle Lys Phe Tyr Thr Lys Gln lle Leu Glu Gly Leu Lys Tyr Leu His
                    230
                                         235
                                                             240
225
```

Glu Asn Gln lle Val His Arg Asp lle Lys Gly Asp Asn Val Leu Val

<212> PRT

Asn	Thr	Tyr	Ser	G1v	Val	Val	Lvs	He	Ser	Asn	Leu	Glv	Thr	Ser	Lvs
ASII	1111	1 7 1	260	dly	, 41	101	Lys	265	501	пор	Lou	01,	270	001	2,5
Ara	ىرە ا	Λla		Val	Asn	Pro	Cve		Glu	Thr	Phe	Thr		Thr	Leu
m g	Lcu	275	01,	741	11311	110	280	1111	Olu	1111	1 110	285	019		Boa
Gln	Tvr		Ala	Pro	Glu	11e		Asn	Gln	Glv	Pro		Glv	Tvr	G1 v
OIII	290	nic c	mia	110	GTU	295	110	пор	OIN	01)	300	8	01,	. y 1	017
Ala		Ala	Asn	Tle	Trp		Leu	Glv	Cvs	Thr		He	Gln	Met	Ala
305	110	mu	пор	110	310	001	Lea	01)	0,0	315	110	110	014	.,,,,	320
	Ser	Lvs	Pro	Pro	Phe	His	Glu	Leu	G1 v		Pro	Gln	Ala	Ala	
		-,-		325					330					335	
Phe	Lys	Val	Gly		Phe	Lys	Ile	His		Glu	Ile	Pro	Glu		Leu
			340					345					350		
Ser	Ala	Glu	Ala	Arg	Ala	Phe	lle	Leu	Ser	Cys	Phe	Glu	Pro	Asp	Pro
		355					360					365			
His	Lys	Arg	Ala	Thr	Thr	Ala	Glu	Leu	Leu	Arg	Glu	G1 y	Phe	Leu	Arg
	370					375					380				
Gln	Val	Asn	Lys	Gly	Lys	Lys	Asn	Arg	Ile	Ala	Phe	Lys	Pro	Ser	Glu
385					390					395					400
Gly	Pro	Arg	Gly	Val	Val	Leu	Ala	Leu	Pro	Thr	Gln	Gly	Glu	Pro	Met
				405					410					415	
Ala	Thr	Ser	Ser	Ser	Glu	His	Gly	Ser	Va]	Ser	Pro	Asp	Ser	Asp	Ala
			420					425					430		
Gln	Pro	Asp	Ala	Leu	Phe	Glu	Arg	Thr	Arg	Ala	Pro	Arg	His	His	Leu
		435					440					445			
Gly	His	Leu	Leu	Ser	Val		Asp	Glu	Ser	Ser		Leu	G]u	Asp	Arg
	450				_	455					460				
	Leu	Ala	Ser	Ser	Pro	Glu	Asp	Arg	Asp		Gly	Leu	Phe	Leu	
465	,		0	0.1	470		. 1	7.1		475 T		7.7		T	480
Arg	Lys	Asp	Ser		Arg	Arg	Ala	11e		lyr	Lys	11e	Leu		GIU
C1	Cln	Aan	Cla	485 Vo.1	11 0	Com	Aan	Lou	490	Clu	Cva	Vol.	A1 0	495	Son
GIU	6111	ASII	500	val	Ala	sei	ASH	505	GIII	Olu	Cys	vai	510	0111	261
Sor	C111	Clu		Hic	Leu	Sor	Val		Hic	م۱۱	Lvc	Gln		ماا	Glv
261	oru	515	Leu	1112	Leu	261	520	Oly	1113	116	ris	525	116	116	Oly
He	Leu		Asn	Phe	Пе	Arø		Pro	Glu	His	Arø		Met	Ala	Thr
110	530	8			210	535	-01				540				

Thr	He	Ser	Lys	Leu	Lys	Val	Asp	Leu	Asp	Phe	Asp	Ser	Ser	Ser	lle
545					550					555					560
Ser	Gln	Ile	His	Leu	Val	Leu	Phe	Gly	Phe	Gln	Asp	Ala	Val	Asn	Lys
				565					570					575	
He	Leu	Arg	Asn	His	Leu	He	Arg	Pro	His	Trp	Met	Phe	Ala	Met	Лsp
			580					585					590		
Asn	He	Ile	Arg	Arg	Ala	Val	Gln	Ala	Ala	Val	Thr	11e	Leu	Ile	Pro
		595					600					605			
Glu	Leu	Arg	Ala	His	Phe	Glu	Pro	Thr	Cys	Glu	Thr	Glu	Gly	Val	Asp
	610					615					620				
Lys	Asp	Met	Asp	Glu	Ala	Glu	Glu	Gly	Tyr	Pro	Pro	Ala	Thr	Gly	Pro
625					630					635					640
Gly	Gln	Glu	Ala	Gln	Pro	His	Gln	Gln	His	Leu	Ser	Leu	Gln	Leu	Gly
				645					650					655	
Glu	Leu	Arg	Gln	Glu	Thr	Asn	Arg	Leu	Leu	Glu	His	Leu	Val	Glu	Lys
			660					665					670		
Glu	Arg	Glu	Tyr	Gln	Asn	Leu	Leu	Arg	Gln	Thr	Leu	Glu	Gln	Lys	Thr
		675					680					685			
Gln	Glu	Leu	Tyr	His	Leu	Gln	Leu	Lys	Leu	Lys	Ser	Asn	Cys	Ile	Thr
	690					695					700				
Glu	Asn	Pro	Ala	Gly	Pro	Tyr	Gly	Gln	Arg	Thr	Asp	Lys	Glu	Leu	He
705					710					715					720
Asp	Trp	Leu	Arg	Leu	Gln	Gly	Ala	Asp	Ala	Lys	Thr	lle	Glu	Lys	lle
				725					730					735	
Val	Glu	Glu	Gly	Tyr	Thr	Leu	Ser	Asp	lle	Leu	Asn	Glu	lle	Thr	Lys
			740					745					750		
Glu	Asp	Leu	Arg	Tyr	Leu	Arg	Leu	Arg	Gly	Gly	Leu	Leu	Cys	Arg	Leu
		755					760					765			
Trp	Ser	Ala	Val	Ser	Gln	Tyr	Arg	Arg	Ala	Gln	Glu	Ala	Ser	Glu	Thr
	770					775					780				
Lys	Asp	Lys	Ala												
785															

<211> 266

<212	2> PF	RT													
<213	3> Ho	omo s	sapie	ens											
<400)> 55	56													
Met	Leu	lle	Asn	Val	He	Ser	Gly	Ser	Ser	Tyr	Met	He	His	Ser	Thr
1				5					10					15	
Gln	Leu	He	Glu	Asn	Ala	Glu	Leu	Arg	Phe	His	Thr	Asp	Glu	Gln	Leu
			20					25					30		
Met	Thr	Leu	Phe	Met	Gln	Leu	Gln	Thr	Ala	Val	Arg	Ser	Arg	Met	His
		35					40					45			
Pro	Phe	Tyr	lle	Thr	His	lle	Arg	Ala	His	Thr	Pro	Leu	Pro	Gly	Pro
	50					55					60				
Leu	Thr	Ala	Gly	Asn	Gln	Met	Ala	Asp	Arg	Leu	Val	Ala	Thr	Ala	11e
65					70					75					80
Ser	Asn	Ala	Arg	His	Phe	His	Asn	Leu	Thr	Arg	Val	Asn	Ala	Ser	G1y
				85					90					95	
Leu	Lys	Arg	Arg	Tyr	Ser	Ser	Thr	Arg	Lys	Glu	Ala	Lys	Ala	lle	He
			100					105					110		
Gln	Arg	Cys	Pro	Thr	Cys	Gln	Met	Val	His	Ser	Ser	Ser	Phe	Thr	G1 y
		115					120					125			
Gly	Val	Asn	Pro	Arg	Arg	Leu	Glu	Pro	Asn	Ser	Leu	Trp	Glu	Met	Asp
	130					135					140				
Val	Thr	His	Va]	Pro	Ser	Phe	Gly	Arg	Leu	Ala	Tyr	Val	His	Ala	Cys
145					150					155					160
Val	Asp	Thr	Phe	Ser	Leu	Trp	Ala	Ala	Cys	Gln	Ser	Gly	Glu	Ser	Ser
				165					170					175	
Ala	Tyr	Val	Lys	Arg	His	Leu	Leu	Gln	Cys	Phe	Val	Val	lle	Gly	11e
			180					185					190		
Leu	Ala	Ser	He	Lys	Thr	Asp	Asn	Ala	Pro	Gly	Tyr	Thr	Ser	Gln	Ala
		195					200					205			
Leu	Ala	Thr	Phe	Phe	Ser	lle	Arg	Asn	He	Lys	His	He	Thr	Gly	He
	210					215					220				
	Tyr	Asn	Ser	Gln		Gln	Ala	lle	Val		Arg	Met	Asn	Leu	
225					230					235					240

Leu Lys Gln Gln Leu Gln Lys Gln Lys Gly Glu Asn Arg Asp Tyr Gly

Thr Pro His Met Gln Leu Asn Arg lle Ile
260 265

<210> 4934 <211> 279 <212> PRT <213> Homo sapiens <400> 557 Met Gly Thr Ser Cys Leu Pro Asp Thr Phe Thr Lys Leu Ile Asn Pro Gln Glu Asn Thr Cys Ser Leu Glu Glu Phe Val Leu Gln Leu Glu Leu Ser Gly Tyr Ser Pro Glu Asp Leu Thr Ala Ala Leu Glu lle Leu Glu Ala Ile 11e Ala Thr Gly Cys Phe Gly 11e Asp Lys Glu Glu Leu Arg Arg Arg Phe Ser Ala Leu Glu Lys Ala Gly Gly Gly Arg Thr Arg Thr Phe Ala Asp Cys Ile Gln Ala Leu Leu Glu Gln His Gln Val Leu Glu Val Gly Gly Asn Thr Ala Arg Leu Val Ala Met Gly Ser Ala Trp Pro Trp Leu Leu His Ser Val Arg Leu Lys Asp Arg Glu Asp Ala Asp Ile Gln Arg Glu Asp Pro Gln Ala Arg Pro Leu Glu Gly Ser Ser Ser Glu Asp Ser Pro Pro Glu Gly Gln Ala Pro Pro Ser His Ser Pro Arg Gly Thr Lys Arg Arg Ala Ser Trp Ala Ser Glu Asn Gly Glu Thr Asp Ala

Glu Gly Thr Gln Met Thr Pro Ala Lys Arg Pro Ala Leu Gln Asp Ser

Asn Leu Ala Pro Ser Leu Gly Pro Gly Ala Glu Asp Gly Ala Glu Ala

Gln Ala Pro Ser Pro Pro Pro Ala Leu Glu Asp Thr Ala Ala Ala Gly 215 220 Ala Ala Gln Glu Asp Gln Glu Gly Val Gly Glu Phe Ser Ser Pro Gly 230 235 240 225 Gln Glu Gln Leu Ser Gly Gln Ala Gln Pro Pro Glu Gly Ser Glu Asp 250 245 Pro Arg Gly Thr Ala Arg Leu Val Pro His Pro Thr Ser Pro His Pro 260 270 265 Gly Phe Pro Ser Pro Pro Pro 275

<210> 4935

<211> 325

<212> PRT

<213> Homo sapiens

<400> 558

Met Pro Ser Arg 11e Leu Asp Lys Glu Leu Leu Ser Gly 11e Pro Asp 5 10 Thr Glu Arg Leu Ser Glu Val Val Asp Asn Arg Ala Pro Gln Arg Asp 20 25 Pro Thr His Arg Ala Pro Gln Arg Asp Pro Thr His Arg Ala Pro Gln 45 Arg Asp Pro Thr His Arg Ala Pro Gln Arg Asp Pro Thr His Arg Ala 55 Pro Gln Arg Asp Pro Thr His Arg Ala Pro Gln Arg Asp Pro Thr His 70 75 80 65 Arg Pro Pro Gln Arg Asp Pro Thr His Arg Ala Pro Gln Arg Asp Pro 85 90 Thr His Arg Ala Pro Gln Arg Asp Pro Thr His Arg Ala Pro Gln Arg 105 Gly Pro Arg His Arg Ala Pro Gln Arg Ser Pro Arg His Arg Ala Pro 115 125 Gln Arg Asp Pro Thr His Arg Ala Pro Gln Arg Gly Pro Arg His Arg

Ala	Pro	Gln	Arg	Λsp	Pro	Thr	His	Arg	Ala	Pro	Gln	Arg	Gly	Pro	Arg
145					150					155					160
His	Arg	Ala	Pro	Gln	Arg	Gly	${\tt Pro}$	Thr	His	Arg	Ala	Pro	Gln	Arg	Gly
				165					170					175	
Pro	Arg	His	Arg	Ala	Pro	Gln	Arg	Gly	Pro	Arg	His	Arg	Ala	Pro	Gln
			180					185					190		
Arg	Gly	Pro	Thr	His	Arg	Ala	Pro	Gln	Arg	Asp	Pro	Gly	Trp	Arg	Ala
		195					200					205			
Pro	Gln	Arg	Gly	Pro	Thr	His	Arg	Ala	Pro	Gln	Arg	lle	Leu	Asp	Ala
	210					215					220				
Glu	Pro	Val	Ser	Gly	Phe	Leu	Asp	Ala	Glu	Ser	Leu	Ser	Gly	He	Leu
225					230					235					240
Val	Val	Glu	Arg	Leu	Ser	Gly	Пе	Leu	Asp	Ala	Glu	Arg	Leu	Ser	G1y
				245					250					255	
He	Leu	Asp	Thr	Glu	Arg	Leu	Ser	Leu	Val	Leu	His	Thr	Glu	Arg	Leu
			260					265					270		
Ser	Gly	Ile	Leu	Val	Gly	Glu	Arg	Leu	Ser	Gly	Va]	Leu	His	Thr	Glu
		275					280					285			
Arg	Leu	Ser	Gly	He	Leu	Val	Gly	Glu	Arg	Leu	Ser	G1 y	lle	Leu	Asp
	290					295					300				
Thr	Glu	Arg	Leu	Ser	Gly	He	Leu	Asp	Thr	Glu	Pro	Leu	Ser	Glu	
305					310					315					320
Gln	He	llis	Ser												
				325											
401	n\ **	20.6													
	0> 49														
	1> 37														
<21:	/	١ I													
101	3> Ho														

Met Met Leu Ser Arg Lys Leu Pro Val His Val Asp Asp Pro Leu Ser

Thr Thr Leu Phe Ser Ser Gln Leu Asn Arg Arg 11e Ser Asn Met Asp

<400> 559

			20					25					30		
Asp	Lys	Val	Tyr	Lys	Met	Ser	Arg	Ala	Leu	Ala	Glu	lle	Lys	Lys	Arg
		35					40					45			
Phe	Gln	Lys	Thr	Val	Thr	Gln	Phe	He	Asn	Ser	He	Leu	Leu	Ala	Ala
	50					55					60				
Gly	Leu	Phe	Thr	lle	Glu	Tyr	Pro	Thr	Lys	Lys	Glu	Glu	Glu	Glu	Phe
65					70					75					80
Val	Arg	Phe	Lys	Met	Arg	Ser	Arg	Thr	His	Pro	Glu	Arg	Leu	${\tt Pro}$	Lys
				85					90					95	
Leu	Ser	Leu	Tyr	Ser	Gly	Glu	Ser	Leu	Leu	Arg	Ser	Gln	Ser	Gly	His
			100					105					110		
Leu	Glu	Ser	Ser	He	Ala	Glu	Thr	Leu	Lys	Asp	Glu	Pro	Glu	Ser	Ala
		115					120					125			
Pro	Val	Ser	Pro	Val	Arg	Lys	Thr	Thr	Lys	He	His	Thr	Lys	Ala	Lys
	130					135					140				
Val	Thr	Ser	Arg	Gly	Lys	Ala	Arg	Glu	Gly	Arg	Ser	Pro	Thr	Arg	Trp
145					150					155					160
Ala	Ala	Leu	Pro	Ser	Asp	Cys	Pro	Leu	Val	Leu	Arg	Lys	Leu	Met	Leu
				165					170					175	
Lys	Glu	Asp	Thr	Arg	Ala	Gly	Cys	Lys	Cys	Leu	Val	Lys	Ala	Pro	Leu
			180					185					190		
Val	Ser		Val	Glu	Leu	Glu		Phe	Leu	Leu	Ala		Arg	Asp	Pro
		195					200			_		205	_		
Ser		Val	Leu	Val	Phe	Gly	Пе	He	Ser	Ser		Asn	Tyr	Thr	Ser
æ.	210	0.1	,	0.1	m	215	,		m		220			٥,١	0.1
	Gly	GIn	Leu	GIn		Leu	Leu	Asn	lhr		lyr	Asn	His	GIn	
225	C1		0.1	C	230		7.1	C1	C	235	т		C	Tr.	240
Arg	GIŸ	Arg	Gly		Pro	Cys	11e	GIn		Arg	lyr	Asp	Ser		Arg
1	Lan	C1	Т	245	1	A	C	Daga	250	C1	C1	Α	D	255 P	1
Leu	Leu	GIN	260	ASP	Leu	Asp	ser		Leu	GIII	GJu	ASP		Pro	Leu
Mot	Vol	Lvc		Acn	Son	Vol	Vo.1	265	Cly	Mot	11.5	Lou	270 Mot	Dho	Alo
Met	val	275	rys	ASII	261	Val	280	0111	01 y	Me t	116	285	me t	rne	MIa
Glv	Ġlv		الم أ	He	Pho	Gly		Arg	Val	يرم ا	Aen		Tur	Glv	اما
51 y	290	r y o	Leu	110	1 116	295	ОТУ	m g	101	Leu	300	01 y	ı yı	01 y	ren
Ser		Gln	Asn	Leu	Leu	Lvs	Gln	Πe	Phe	Arg		Gln	Gln	Asp	Tvr

Lys Met Gly Tyr Phe Leu Pro Asp Asp Tyr Lys Phe Ser Val Pro Asn Ser Val Leu Ser Leu Glu Asp Ser Glu Ser Val Lys Lys Ala Glu Ser Glu Asp lle Gln Gly Ser Ser Ser Ser Leu Ala Leu Glu Asp Tyr Val Glu Lys Glu Leu Ser Leu Glu Ala Glu Lys

<210> 4937

<211> 455

<212> PRT

<213> Homo sapiens

<400> 560 Met Gly Val Ile Ile Ser Gly Ser Leu Pro Pro Phe Ser Ser Pro Leu Gln Val Gly Pro Val Ser Leu Leu Ala Ile Gly Val Leu Thr Val His Cys Met Val Ile Leu Leu Asn Cys Ala Gln His Leu Ser Gln Arg Leu Gln Lys Thr Phe Val Asn Tyr Gly Glu Ala Thr Met Tyr Gly Leu Glu Thr Cys Pro Asn Thr Trp Leu Arg Ala His Ala Val Trp Gly Arg Trp Asn Leu Ala Leu Ser Pro Arg Leu Glu Cys Ser Gly Lys 11e Ser Ala His Cys Asn Pro His Leu Gln Gly Ser Ser Asn Ser Pro Ala Gln Ala Ser Arg Val Ala Gly 11e Tyr Arg Tyr Thr Val Ser Phe Leu Leu Val

The Thr Gln Leu Gly Phe Cys Ser Val Tyr Phe Met Phe Met Ala Asp

Asn Leu Gln Gln Met Val Glu Lys Ala His Val Thr Ser Asn Ile Cys

145					150					155					160
Gln	Pro	Arg	Glu	Ile	Leu	Thr	Leu	Thr	Pro	Ile	Leu	Asp	He	Arg	Phe
				165					170					175	
Tyr	Met	Leu	He	He	Leu	Pro	Phe	Leu	He	Leu	Leu	Val	Phe	lle	Gln
			180					185					190		
Asn	Leu	Lys	Val	Leu	Ser	Val	Phe	Ser	Thr	Leu	Ala	Asn	lle	Thr	Thr
		195					200					205			
Leu	Gly	Ser	Met	Ala	Leu	Ile	Phe	Glu	Tyr	Ile	Met	Glu	Gly	lle	Pro
	210					215					220				
Tyr	Pro	Ser	Asn	Leu	Pro	Leu	Met	Ala	Asn	Trp	Lys	Thr	Phe	Leu	Leu
225					230					235					240
Phe	Phe	Gly	Thr	Ala	11e	Phe	Thr	Phe	Glu	Gly	Val	Gly	Met	Val	Leu
				245					250					255	
Pro	Leu	Lys	Asn	Gln	Met	Lys	His	Pro	Gln	Gln	Phe	Ser	Phe	Val	Leu
			260					265					270		
Tyr	Leu	Gly	Met	Ser	Ile	Val	lle	lle	Leu	Tyr	Ile	Leu	Leu	Gly	Thr
		275					280					285			
Leu	Gly	Tyr	Met	Lys	Phe	Gly	Ser	Asp	Thr	Gln	Ala	Ser	Ile	Thr	Leu
	290					295					300				
Asn	Leu	Pro	Asn	Cys	Trp	Leu	Tyr	Gln	Ser	Val	Lys	Leu	Met	Tyr	Ser
305					310					315					320
He	G1 y	He	Phe		Thr	Tyr	Ala	Leu		Phe	His	Val	Pro	Ala	Glu
				325					330					335	
He	He	He		Phe	Ala	He	Ser		Val	Ser	Glu	Ser		Ala	Leu
			340	_				345					350		
Phe	Val		Leu	Ser	Val	Arg		Ala	Leu	Val	Cys		Thr	Cys	Val
0		355		7.7			360		_			365			
Ser		He	Leu	He	Pro		Leu	Asp	Leu	Val		Ser	Leu	Val	G] y
C	370	C .	C .	C		375					380				0.1
	vai	ser	Ser	Ser		Leu	Ala	Leu	11e		Pro	Ala	Leu	Leu	
385	W - 1	ті.	DL -	т	390	C1		he .	C	395	W 1	TI		4.1	400
116	val	116	rne		ser	UIU	Asp	Met		uys	val	ınr	116	Ala	Lys
Acn	110	Mo+	11.	405	11.	Ve 1	C1	ىنم ا	410	C1	Cv	11-	Dk -	415	Tl
rəp	116	Met	420	261	116	va1			Leu		Cys	116	430	Gly	1111

Tyr Gln Ala Leu Tyr Glu Leu Pro Gln Pro Ile Ser His Ser Met Ala
435

Asn Ser Thr Gly Val His Ala
450

455

<210> 4938

<211> 162

<212> PRT

<213> Homo sapiens

<400> 561

Met Ser Ile Thr Arg Leu His Glu Gln Lys Leu Val Gln His Val Val

1 5 10 15

Ser Gln Asn Cys Asp Gly Leu His Leu Arg Ser Gly Leu Pro Arg Thr
20 25 30

Ala Ile Ser Glu Leu His Gly Asn Met Tyr Ile Glu Val Ser Ser Pro 35 40 45

Ala Gly Thr Gln Gly Leu His Gly Gln Ala Gly Pro Thr His Cys Ala 50 55 60

Leu Leu Pro Leu Gly Leu Tyr Leu Leu Arg Ser Gln Gln Gly Val Arg
65 70 75 80

Ala Gly Val Arg Cys Asp Gly Ala His Cys Pro Pro Gln Thr Pro Asp
85 90 95

Arg Pro Asp Leu Pro Gln Val Trp Asp Pro Ala Ala Gly His His Cys 100 105 110

Ala Leu Trp Gly Glu Gly Asp Val Gly Ala Ala Ser Glu Leu Gly Ser 115 120 125

Gly Asp Arg Gly Cys Gln Gln Ser Arg His His Pro Val Ser Arg Val 130 135 140

Gln Pro Glu Gly Ser Lys Glu Val Pro Thr Pro Leu Val His Asp Gln 145 150 155 160

Ala Pro

```
<210> 4939
<211> 567
<212> PRT
<213> Homo sapiens
<400> 562
Met Asp Asp Lys Asp lle Asp Lys Glu Leu Arg Gln Lys Leu Asn Phe
 1
                                      10
Ser Tyr Cys Glu Glu Thr Glu Ile Glu Gly Gln Lys Lys Val Glu Glu
             20
                                 25
Ser Arg Glu Ala Ser Ser Gln Thr Pro Glu Lys Gly Glu Val Gln Asp
                             40
Ser Glu Ala Lys Gly Thr Pro Pro Trp Thr Pro Leu Ser Asn Val His
     50
                         55
                                              60
Glu Leu Asp Thr Ser Ser Glu Lys Asp Lys Glu Ser Pro Asp Gln Ile
                     70
                                          75
Leu Arg Thr Pro Val Ser His Pro Leu Lys Cys Pro Glu Thr Pro Ala
                 85
                                      90
Gln Pro Asp Ser Arg Ser Lys Leu Leu Pro Ser Asp Ser Pro Ser Thr
            100
                                105
                                                     110
Pro Lys Thr Met Leu Ser Arg Leu Val 11e Ser Pro Thr Gly Lys Leu
                            120
                                                 125
Pro Ser Arg Gly Pro Lys His Leu Lys Leu Thr Pro Ala Pro Leu Lys
    130
                        135
                                             140
Asp Glu Met Thr Ser Leu Ala Leu Val Asn Ile Asn Pro Phe Thr Pro
                    150
                                         155
Glu Ser Tyr Lys Lys Leu Phe Leu Gln Ser Gly Gly Lys Arg Lys Ile
                165
                                    170
Arg Gly Asp Leu Glu Glu Ala Gly Pro Glu Glu Gly Lys Gly Gly Leu
            180
                                185
                                                     190
Pro Ala Lys Arg Cys Val Leu Arg Glu Thr Asn Met Ala Ser Arg Tyr
                            200
                                                 205
Glu Lys Glu Phe Leu Glu Val Glu Lys 11e Gly Val Gly Glu Phe Gly
    210
                        215
                                             220
Thr Val Tyr Lys Cys Ile Lys Arg Leu Asp Gly Cys Val Tyr Ala Ile
```

sn Ser Ala

255

ro His Val

77 O

le Ile Gln

∜œr Glu Asn

sp Ile Leu

320

er Met Val

335

ys Vál Gln

50

la Asp Trp

ly His Ala

rg Phe Leu

400

ys Ala Asp

415

la Glu Ser

33 ()

ly Asn Phe

wu Leu Lys

la Ala Leu

480

lu Glu Leu

495

💇 Glu Arg

HO

yr Thr His

<210> 4940

<211> 140

<212> PRT

<213> Homo sapiens

<400> 563

Met Asp Phe lle Val Ala Ala Ser Asn Leu Arg Ala Glu Asn Tyr Asp

1 5 10 15

Ile Pro Ser Ala Asp Arg His Lys Ser Lys Leu Ile Ala Gly Lys Ile

20 25 · 30

Ile Pro Ala Ile Ala Thr Thr Thr Ala Ala Val Val Gly Leu Val Cys 35 40 45

Leu Glu Leu Tyr Lys Val Val Gln Gly His Arg Gln Leu Asp Ser Tyr 50 55 60

Lys Asn Gly Phe Leu Asn Leu Ala Leu Pro Phe Phe Gly Phe Ser Glu 65 70 75 80

Pro Leu Ala Ala Pro Arg His Gln Tyr Tyr Asn Gln Glu Trp Thr Leu 85 90 95

Trp Asp Arg Phe Glu Val Gln Gly Leu Gln Pro Asn Gly Glu Glu Met
100 105 110

Thr Leu Lys Gln Phe Leu Asp Tyr Phe Lys Val Arg Pro Leu Pro Tyr 115 120 125

Ser Val Thr Pro Pro Gln Gly Ala Arg Cys Thr Arg 130 135 140

<210> 4941

<211> 477

	<212	2> PF	RT.													
	<213	3> Ho	omo s	sapie	ens											
	<400)> 56	54										•			
	Met	Ser	Ala	Ala	Pro	G1 y	Leu	Leu	His	Gln	Glu	Leu	Ser	Čys	Pro	Leu
	l				5					10					15	
	Cys	Leu	G]n	Leu	Phe	Asp	Ala	Pro	Val	Thr	Ala	Glu	Cys	Gly	His	Ser
				20					25					30		
	Phe	Cys	Arg	Ala	Cys	Leu	Gly	Arg	Val	Ala	Gly	Glu	Pro	Ala	Ala	Asp
•			35					40					45			
	Gly	Thr	Val	Leu	Cys	Pro	Cys	Cys	Gln	Ala	Pro	Thr	Arg	Pro	Gln	Ala
		50					55					60				
	Leu	Ser	Thr	Asn	Leu	Gln	Leu	Ala	Arg	Leu	Val	Glu	Gly	Leu	Ala	Glr
	65					70					75					80
	Val	Pro	Gln	Gly	His	Cys	Glu	Glu	His	Leu	Asp	Pro	Leu	Ser	lle	Tyr
					85					90					95	
	Cys	Glu	Gln	Asp	Arg	Ala	Leu	Val	Cys	Gly	Val	Cys	Ala	Ser	Leu	Gly
				100					105					110		
	Ser	His	Arg	Gly	His	Arg	Leu	Leu	Pro	Ala	Ala	Glu	Ala	His	Ala	Arg
			115					120					125			
	Leu	Lys	Thr	Gln	Leu	Pro	Gln	Gln	Lys	Leu	Gln	Leu	Gln	Glu	Ala	Cys
		130					135					140				
	Met	Arg	Lys	Glu	Lys	Ser	Val	Ala	Val	Leu	Glu	His	Gln	Leu	Val	G1ı
	145					150	,				155					160
	Val	Glu	Glu	Thr	Val	Arg	Gln	Phe	Arg	Gly	Ala	Val	Gly	Glu	Gln	Leı
					165					170					175	
	Gly	Lys	Met	Arg	Val	Phe	Leu	Ala	Ala	Leu	Glu	Gly	Ser	Leu	Asp	Cys
				180					185					190		
	Glu	Ala	Glu	Arg	Val	Arg	Gly	Glu	Ala	Gly	Val	Ala	Leu	Arg	Arg	G1ı
			195					200					205			
	Leu	Gly	Ser	Leu	Asn	Ser	Tyr	Leu	Glu	Gln	Leu	Arg	Gln	Met	Glu	Lys
		210					215					220				
	Val	Leu	Glu	Glu	Va]	Ala	Asp	Lys	Pro	G] n	Thr	Glu	Phe	Leu	Met	Lys
	225					230					235					240

Tyr Cys Leu Val Thr Ser Arg Leu Gln Lys Ile Leu Ala Glu Ser Pro

```
Pro Pro Ala Arg Leu Asp 11e Gln Leu Pro 11e 11e Ser Asp Asp Phe
                                265
Lys Phe Gln Val Trp Arg Lys Met Phe Arg Ala Leu Met Pro Ala Leu
                            280
                                                285
Glu Glu Leu Thr Phe Asp Pro Ser Ser Ala His Pro Ser Leu Val Val
                        295
                                            300
Ser Ser Ser Gly Arg Arg Val Glu Cys Ser Glu Gln Lys Ala Pro Pro
                    310
                                        315
                                                            320
Ala Gly Glu Asp Pro Arg Gln Phe Asp Lys Ala Val Ala Val Val Ala
                                    330
                325
                                                        335
His Gln Gln Leu Ser Glu Gly Glu His Tyr Trp Glu Val Asp Val Gly
            340
                                345
Asp Lys Pro Arg Trp Ala Leu Gly Val lle Ala Ala Glu Ala Pro Arg
        355
                            360
                                                365
Arg Gly Arg Leu His Ala Val Pro Ser Gln Gly Leu Trp Leu Leu Gly
                        375
Leu Arg Glu Gly Lys Ile Leu Glu Ala His Val Glu Ala Lys Glu Pro
                    390
                                        395
Arg Ala Leu Arg Ser Pro Glu Arg Arg Pro Thr Arg 11e Gly Leu Tyr
                405
                                    410
                                                        415
Leu Ser Phe Gly Asp Gly Val Leu Ser Phe Tyr Asp Ala Ser Asp Ala
                                425
Asp Ala Leu Val Pro Leu Phe Ala Phe His Glu Arg Leu Pro Arg Pro
                                                445
        435
                            440
Val Tyr Pro Phe Phe Asp Val Cys Trp His Asp Lys Gly Lys Asn Ala
    450
                        455
Gln Pro Leu Leu Val Gly Pro Glu Gly Ala Glu Ala
                    470
                                        475
```

<211> 505

<212> PRT

<213> Homo sapiens

<400> 565

Met	Ala	Val	Ala	Leu	Asp	Ser	Gln	He	Asp	Ala	Pro	Leu	Glu	Val	Glu
l				5					10					15	
Gly	Cys	Leu	lle	Met	Lys	Val	Glu	Lys	Asp	Pro	Glu	Trp	Ala	Ser	Glu
			20					25					30		
Pro	lle	Leu	Glu	Gly	Ser	Asp	Ser	Ser	Glu	Thr	Phe	Arg	Lys	Cys	Phe
		35					40					45			
Arg	Gln	Phe	Cys	Tyr	Glu	Asp	Val	Thr	Gly	Pro	His	Glu	Ala	Phe	Ser
	50					55					60				
Lys	Leu	Trp	Glu	Leu	Cys	Cys	Arg	Trp	Leu	Lys	Pro	Glu	Met	Arg	Ser
65					70					75					80
Lys	Glu	Gln	Ile	Leu	Glu	Leu	Leu	Val	Ile	Glu	Gln	Phe	Leu	Thr	Ile
				85					90					95	
Leu	Pro	${\tt Glu}$	Lys	He	Gln	Ala	Trp	Ala	Gln	Lys	Gln	Cys	Pro	Gln	Ser
			100					105					110		
Gly	Glu	Glu	Ala	Val	Ala	Leu	Val	Val	His	Leu	Glu	Lys	Glu	Thr	Gly
		115					120					125			
Arg	Leu	Arg	Gln	Gln	Val	Ser	Ser	Pro	Val	His	Arg	Glu	Lys	His	Ser
	130					135					140				
Pro	Leu	Gly	Ala	Ala	Trp	Glu	Val	Ala	Asp	Phe	Gln	Pro	Glu	Gln	Val
145					150					155					160
Glu	Thr	Gln	Pro	Arg	Ala	Val	Ser	Arg	Glu	Glu	Pro	Gly	Ser	Leu	His
				165					170					175	
Ser	Gly	His	Gln	Glu	Gln	Leu	Asn	Arg	Lys	Arg	Glu	Arg	Arg	Pro	Leu
			180					185					190		
Pro	Lys	Asn	Ala	Arg	Pro	Ser	Pro	Trp	Va]	Pro	Ala	Leu	Ala	Asp	Glu
		195					200					205			
Trp	Asn	Thr	Leu	Asp	Gln	Glu	Val	Thr	Thr	Thr	Arg	Leu	Pro	Ala	Gly
	210					215					220				
Ser	Gln	Glu	Pro	Val		Asp	Val	His	Val		Arg	Gly	Phe	Ser	
225					230					235					240
Arg	Lys	Ser	Val		Gln	He	Pro	Ala	Gln	Arg	Asp	Leu	Tyr	Arg	Asp
				245					250					255	
Phe	Arg	Lys		Asn	Val	GI y	Asn		Va]	Ser	Leu	Gly		Ala	Val
			260					265					270		
Ser	Thr		Asn	Lys	He	Thr		Leu	Glu	Gln	Arg		Glu	Pro	Trp
		275					280					285			

Thr	Leu	Gly	Leu	His	Ser	Ser	Asn	Lys	Arg	Ser	lle	Leu	Arg	Ser	Asn
	290					295					300				
Tyr	Val	Lys	Glu	Lys	Ser	Val	His	Ala	Πe	Gln	Val	Pro	Ala	Arg	Ser
305					310					315					320
Ala	Gly	Lys	Thr	Trp	Arg	Glu	Gln	Gln	Gln	Trp	G1 y	Leu	Glu	Лsp	Glu
				325					330					335	
Lys	Пе	Ala	Gly	Val	His	Trp	Ser	Tyr	Glu	Glu	Thr	Lys	Thr	Phe	Leu
			340					345					350		
Ala	He	Leu	Lys	Glu	Ser	Arg	Phe	Tyr	Glu	Thr	Leu	Gln	Ala	Cys	Pro
		355					360					365			
Arg	Asn	Ser	Gln	Val	Tyr	Gly	Ala	Val	Ala	Glu	Trp	Leu	Arg	Glu	Cys
	370					375					380				
Gly	Phe	Leu	Arg	Thr	Pro	Glu	Gln	Cys	Arg	Thr	Lys	Phe	Lys	Ser	Leu
385					390					395					400
Gln	Lys	Ser	Tyr	Arg	Lys	Val	Arg	Asn	Gly	His	Met	Leu	Glu	Pro	Cys
				405					410					415	
Ala	Phe	Phe	Glu	Asp	Met	Asp	Ala	Leu	Leu	Asn	Pro	Ala	Ala	Arg	Ala
			420					425					430		
Pro	Ser	Thr	Asp	Lys	Pro	Lys	Glu	Met	Ile	Pro	Val	Pro	Arg	Leu	Lys
		435					440					445			
Arg	11e	Ala	He	Ser	Ala	Lys	Glu	His	lle	Ser	Leu	Val	Glu	Glu	Glu
	450					455					460				
Glu	Ala	Ala	Glu	Asp	Ser	Asp	Asp	Asp	Glu	He	Gly	lle	Glu	Phe	He
465					470					475					480
Arg	Lys	Ser	Glu	He	His	Gly	Ala	Pro	Va]	Leu	Phe	Gln	Asn	Leu	Ser
				485					490					495	
Gly	Lys	Asn	Cys	Ala	Leu	Phe	Leu	Trp							
			500					505							

<211> 133

<212> PRT

<213≻ Homo sapiens

<400> 566

Met Pro Pro Leu Ala Arg Ser Pro Arg Ala Ser lle Val Ala Leu Arg 10 Lys Met Ile Thr Ser Phe Leu Ala Leu Trp Trp His Val Cys Val Cys 20 25 30 lle Cys lle Tyr Thr His lle Tyr lle Asn Thr Gln Thr His lle Tyr Thr Asn Thr Arg lle His Phe Arg Pro Gln Phe Leu Ala His Asn Ser 55 His Ser Pro Cys Tyr Thr Leu Leu Leu Gln His Trp Val Cys Gln Ala 70 75 Ser Gly Asp Asn His Ser Asn Leu Leu Pro Phe Leu Leu Pro Ala Gln 90 85 Asp Arg Thr Leu Ile Phe Pro Thr Phe Leu Met Val Gly His Lys Thr 100 105 110 His Ser Arg Asp Gly Pro Thr Pro Tyr Pro Ala Arg Arg Asn Ala Ala 120 115 125 Val Met Lys Leu Pro 130

<210> 4944 <211> 502

<212> PRT

<213> Homo sapiens

<400> 567

Met Val Gln Tyr His Gly Ala Glu Ala Ala Gln Arg Phe lle Leu Thr 1 5 10 15

Val Met Asn Met Val Tyr Asn Met Phe Gln His Gln Ser Leu Gly 11e 20 25 30

Lys 11e Asn 11e Gln Val Thr Lys Leu Val Leu Leu Arg Gln Arg Pro 35 40 45

Ala Lys Leu Ser lle Gly His His Gly Glu Arg Ser Leu Glu Ser Phe 50 55 60

Cys His Trp Gln Asn Glu Glu Tyr Gly Gly Ala Arg Tyr Leu Gly Asn 65 70 75 80

Asn	Gln	Val	Pro	Gly	Gly	Lys	Asp	Asp		Pro	Leu	Val	Asp		Ala
				85					90					95	
Val	Phe	Val	Thr	Arg	Thr	Asp	Phe	Cys	Val	His	Lys	Asp	Glu	Pro	Cys
			100					105					110		
Asp	Thr	Val	Gly	He	Ala	Tyr	Leu	Gly	Gly	Val	Cys	Ser	Ala	Lys	Arg
		115					120	•				125			
Lys	Cys	Val	Leu	Ala	Glu	Asp	Asn	Gly	Leu	Asn	Leu	Ala	Phe	Thr	Ile
	130					135					140				
Ala	His	Glu	Leu	Gly	His	Asn	Leu	Gly	Met	Asn	His	Asp	Asp	Asp	His
145					150					155					160
Ser	Ser	Cys	Ala	Gly	Arg	Ser	His	lle	Met	Ser	Gly	Glu	Trp	Val	Lys
				165					170					175	
Gly	Arg	Asn	Pro	Ser	Asp	Leu	Ser	Trp	Ser	Ser	Cys	Ser	Λrg	Asp	Asp
			180					185			•		190		
Leu	Glu	Asn	Phe	Leu	Lys	Ser	Lys	Val	Ser	Thr	Cys	Leu	Leu	Val	Thr
		195					200					205			
Asp	Pro	Arg	Ser	Gln	His	Thr	Val	Arg	Leu	Pro	His	Lys	Leu	Pro	Gly
	210					215					220				
Met	His	Tyr	Ser	Ala	Asn	Glu	Gln	Cys	Gln	lle	Leu	Phe	Gly	Met	Asn
225					230					235					240
Ala	Thr	Phe	Cys	Arg	Asn	Met	Glu	His	Leu	Met	Cys	Ala	Gly	Leu	Trp
				245					250					255	
Cys	Leu	Val	Glu	Gly	Asp	Thr	Ser	Cys	Lys	Thr	Lys	Leu	Asp	Pro	Pro
			260					265					270		
Leu	Asp	Gly	Thr	Glu	Cys	Gly	Ala	Asp	Lys	Trp	Cys	Arg	Ala	G1 y	Glu
		275					280					285			
Cys	Val	Ser	Lys	Thr	Pro	lle	Pro	Glu	His	Val	Asp	Gly	Asp	Trp	Ser
	290					295					300				
Pro	Trp	Gly	Ala	Trp	Ser	Met	Cys	Ser	Arg	Thr	Cys	Gly	Thr	Gly	Ala
305					310					315					320
Arg	Phe	Arg	Gln	Arg	Lys	Cys	Asp	Asn	Pro	Pro	Pro	Gly	Pro	Gly	Gly
				325					330					335	
Thr	His	Cys	Pro	Gly	Ala	Ser	Val	Glu	His	Ala	Val	Cys	Glu	Asn	Leu
			340	-				345					350		
Pro	Cys	Pro	Lys	Gl y	Leu	Pro	Ser		Arg	Asp	Gln	Gln	Cys	Gln	Ala
	*	355		-			360					365			

His Asp Arg Leu Ser Pro Lys Lys Gly Leu Leu Thr Ala Val Val Val Asp Asp Lys Pro Cys Glu Leu Tyr Cys Ser Pro Leu Gly Lys Glu Ser Pro Leu Leu Val Ala Asp Arg Val Leu Asp Gly Thr Pro Cys Gly Pro Tyr Glu Thr Asp Leu Cys Val His Gly Lys Cys Gln Lys Ile Gly Cys Asp Gly Ile Ile Gly Ser Ala Ala Lys Glu Asp Arg Cys Gly Val Cys Ser Gly Asp Gly Lys Thr Cys His Leu Ala Lys Gly Asp Phe Ser His Ala Arg Gly Thr Gly Tyr lle Glu Ala Ala Val Ile Pro Ala Gly Ala Arg Arg Ile Arg Val Val Glu Asp Lys Pro Ala His Ser Phe Leu Gly Lys Thr Gln Met Thr

<210> 4945

<211> 356

<212> PRT

<213> Homo sapiens

<400> 568

Arg	Gly	Pro	Leu	Ser	Ser	∙Ala	Pro	Glu	He	Val	His	Glu	Asp	Leu	Lys
				85					90					95	
Met	Gly	Ser	Asp	Gly	G]u	Ser	Asp	Gln	Ala	Ser	Ala	Thr	Ser	Ser	Asp
			100					105					110		
Glu	Val	Gln	Ser	Pro	Val	Arg	Val	Arg	Met	Arg	Asn	His	Pro	Pro	Arg
		115					120					125			
Lys	He	Ser	Thr	Glu	Asp	He	Asn	Lys	Arg	Leu	Ser	Leu	Pro	Λla	Asp
	130					135					140				
Ile	Arg	Leu	Pro	Glu	Gly	Tyr	Leu	Glu	Lys	Leu	Thr	Leu	Asn	Ser	Pro
145					150					155					160
He	Phe	Asp	Lys	Pro	Leu	Ser	Arg	Arg	Leu	Arg	Arg	Va]	Ser	Leu	Ser
				165					170					175	
Glu	He	Gly	Phe	Gly	Lys	Leu	Glu	Thr	Tyr	He	Lys	Leu	Asp	Lys	Leu
			180					185					190		
Gly	Glu	Gly	Thr	Tyr	Ala	Thr	Val	Tyr	Lys	Gly	Lys	Ser	Lys	Leu	Thr
		195					200					205			
Asp	Asn	Leu	Val	Ala	Leu	Lys	Glu	lle	Arg	Leu	Glu	His	Glu	Glu	Gly
	210					215					220				
Ala	Pro	Cys	Thr	Ala		Arg	Glu	Val	Ser		Leu	Lys	Asp	Leu	
225					230					235					240
His	Ala	Asn	He		Thr	Leu	His	Asp		He	His	Thr	Glu		Ser
				245					250					255	
Leu	Thr	Leu		Phe	Glu	Tyr	Leu	Asp	Lys	Asp	Leu	Lys		Tyr	Leu
			260					265					270		
Asp	Asp	-	Gly	Asn	He	He		Met	His	Asn	Val		Val	Gly	Va]
		275					280		_			285			
Gly		Glu	Ala	Gly	Ala		Gly	Gly	Pro	His		Pro	Thr	Pro	Thr
	290	_				295					300				
	Lys	Ser	Pro	Arg		Gly	Leu	Phe	Pro		Ala	Phe	Phe	Ala	
305					310	0.1				315	,	0			320
Ser	Pro	Trp	Arg		Leu	Gly	Pro	Cys		Leu	Leu	Cys	Asp		Ala
				325					330					335	
Leu	Gly	Leu		Ser	Val	Phe	Gly	Arg	Gly	Ala	Val	Pro		61 y	Gly
		C	340					345					350		
Arg	Ala	Ser	Gly												
		355													

<210> 4946 <211> 126 <212> PRT <213> Homo sapiens <400> 569 Met Met Ile Thr Ala His Tyr Ser Leu Asp Phe Thr Gly Ser Gly Asp 10 Phe Pro Thr Ser Ala Ser Arg Val Ala Gly Thr Thr Gly Thr His His 25 His Thr Gln Leu lle Phe Cys Ile Phe Ser Arg Gly Arg Val Leu Pro 35 40 45 Cys Cys Pro Gly Trp Ser Arg Thr Pro Gly Leu Lys Gln Ser Ala His 55 Leu Gly Leu Pro Lys Cys Trp Ile Thr Gly Met Ser Arg Cys Ala Gln 70 75 65 80 Pro Lys Ile Ile Phe Ile Leu Phe Ile Gly Ser Ser Phe Ile Ala Leu 85 90 Glu Ile Gln Ala Ala Arg Tyr Cys Asp Trp Cys lle Trp Thr Tyr Leu 100 105 Trp Ser Lys Thr Gly Thr His Gln Ile Val Ala Leu Ala Asn 120 115 125

<210> 4947

<211> 141

<212> PRT

<213> Homo sapiens

<400> 570

Met Leu Leu Trp Thr Leu Gln Tyr Lys Cys Leu Thr Lys Ser Leu Leu

1 5 10 15

Leu Ile Leu Trp Gly Ile Tyr Ile Lys Val Glu Leu Leu Asp His Leu

Val Val Leu His Phe Thr Phe Phe Arg Asp Cys His 11e Val Phe His Ser Asp Cys Thr Ile Leu Tyr Ser Leu Arg Gln Tyr Ala Arg Val Leu Ile Ser Pro Tyr Ser His Gln His Leu Phe Ser Val Leu Arg Ile Ile Ala Ile Leu Arg Gly Val Met Trp Tyr Leu Ile Glu Val Ser Ile Cys Ile Ser Leu Met Ile Ser Asp Val Glu Cys Phe Phe Met Tyr Phe Leu Ala Ile Cys Ile Ser Pro Leu Lys Lys Tyr Gln Tyr Gln Val lle Cys Ser Cys Leu Ile Glu Leu Leu Cys Cys Gly Gly Phe Tyr

<210> 4948

<211> 152

<212> PRT

<213> Homo sapiens

<400> 571

Met Gly Phe His Tyr Val Gly Gln Ala Gly Leu Glu Leu Leu Thr Ser Gly Asp Pro Leu Ala Ser Ala Phe Gln Ser Ala Gly Ile Ile Gly Val Ser His Arg Thr Trp Gly Gly Tyr Cys Leu Lys Lys Lys Ser Pro Asp Ser Asp Pro Leu Ser Thr Trp Arg Thr Ser Thr Gly Arg Lys Gln Met Leu Gln 11e Phe Lys Tyr Pro Asp Gly Phe Gly Ser Gln Gly Glu Arg Asp Leu Thr Ser Val Tyr His Pro Thr Leu Ser Thr Lys Val Thr 11e

Asn Thr Lys Ser Ile Ala Trp Ala Thr Gly Lys Lys Ala Phe Ile Cys

Ile Asn Val Cys Val Tyr Ile His Cys Phe Phe Phe Lys Arg Leu Gly Leu Pro Leu Ser Pro Arg Leu Glu His Ser Gly Thr Ile Ile Ala His Cys Ser Leu Gln Arg Cys Gly <210> 4949 <211> 168 <212> PRT <213> Homo sapiens <400> 572 Met Tyr Pro Ser Asn Lys Lys Lys Lys Val Trp Arg Glu Glu Lys Glu Arg Leu Leu Lys Met Thr Leu Glu Glu Arg Arg Lys Glu Tyr Leu Arg Asp Tyr lle Pro Leu Asn Ser lle Leu Ser Trp Lys Glu Glu Met Lys Gly Lys Gly Gln Asn Asp Glu Glu Asn Ile Gln Glu Thr Ser Gln Val Lys Lys Ser Leu Thr Glu Lys Val Ser Leu Tyr Arg Gly Asp lle Thr Leu Leu Glu Val Asp Ala Ile Val Asn Ala Ala Asn Ala Ser Leu Leu Gly Gly Gly Val Asp Gly Cys Ile His Arg Ala Ala Gly Pro Cys Leu Leu Ala Glu Cys Arg Asn Leu Asn Gly Cys Asp Thr Gly His Ala Lys lle Thr Cys Gly Tyr Asp Leu Pro Ala Lys Tyr Val Ile His Thr Val Gly Pro Ile Ala Arg Ala Ile Leu Met Val Pro Thr Arg Lys Thr

Leu Gln Ile Ala Ile Asn His Leu

165

```
<210> 4950
<211> 151
<212> PRT
<213> Homo sapiens
<400> 573
Met Gly Leu Gly Gly Val Ser Arg Glu Ala Gly Leu Leu Leu Ser
                                      10
Ser Pro Cys Pro Asp Pro Ser Val Cys Leu Ser Asp Lys Pro Val Pro
                                 25
Glu Glu Ser Glu Gly Pro Gly Ser Pro Pro Pro Tyr Lys Met Ile Gln
         35
                             40
                                                  45
Thr Ile Gly Leu Ser Val Gly Ala Ala Val Ala Tyr Ile Ile Ala Val
                         55
Leu Gly Leu Met Phe Tyr Cys Lys Lys Arg Cys Lys Ala Lys Arg Leu
65
                     70
                                         75
                                                              80
Gln Lys Gln Pro Glu Gly Glu Glu Pro Glu Met Glu Cys Leu Asn Gly
                 85
Gly Pro Leu Gln Asn Gly Gln Pro Ser Ala Glu Ile Gln Glu Glu Val
                                105
Ala Leu Thr Ser Leu Gly Ser Gly Pro Ala Ala Thr Asn Lys Arg His
        115
                            120
                                                 125
Ser Thr Ser Asp Lys Met His Phe Pro Arg Ser Ser Leu Gln Pro 11e
                        135
                                            140
Thr Thr Leu Gly Met Leu Pro
```

<210> 4951

145

<211> 182

<212> PRT

<213> Homo sapiens

<400> 574 Met Leu Pro Thr Ala Ala Gly Phe Ser Ile Trp Gly Gln Val Gly Ala 10 15 Ala Arg Glu Ala Pro Arg Cys Gln Thr Lys Ile Ser Ser Cys Ser Cys 20 25 Pro Thr Ser Ser Val Ser Ala Ala His Gly Pro Gly Pro Asn Glu Arg 40 Ala Arg Gly Leu Gly Gly Leu Pro Asp Pro Ala Leu Ser Pro Arg Val 50 55 Pro Phe Gln Gly Tyr Ala Arg Ile Val Phe Ala Ile Ile Ser Phe Tyr 75 70 Phe Met Pro Cys Cys Pro Leu Thr Ala Ser Ser Phe Tyr Leu Leu Ser 85 90 Gly Leu Leu Asp Ala Phe Asp Gly His Ala Ala Arg Ala Leu Asn Gln 100 105 110 Gly Thr Arg Phe Gly Ala Met Leu Asp Met Leu Thr Asp Arg Cys Ser 120 Thr Met Cys Leu Leu Val Asn Leu Ala Leu Leu Tyr Pro Gly Ala Thr 130 135 140 Leu Phe Phe Gln Ile Ser Met Ser Leu Asp Val Ala Ser His Trp Leu 150 155 His Leu His Arg Ser Ala Ala Ile Leu Gly Ala Trp Ala Thr Trp Arg 165 170 His Tyr Ser Gly Val Gly 180 <210> 4952

<211> 258

<212> PRT

<213> Homo sapiens

<400> 575

Met Gly Thr Leu Thr Phe His Leu Lys Ser Ser Phe Pro Gln Val Leu 1 5 10 15 Arg His Val Asn Gly Gln Asp Gln lle Val Pro Gly Leu Tyr Ala Cys

			20					25					30		
Gly	Glu	Λla	Ala	Cys	Ala	Ser	Val	His	Gly	Ala	Asn	Arg	Leu	Gly	Ala
		35					40					45			
Asn	Ser	Leu	Leu	Asp	Leu	Val	Ala	Phe	Gly	Arg	Ala	Cys	Ala	Pro	Ser
	50					55					60				
He	Glu	Glu	Ser	Cys	Arg	Pro	Gly	Asp	Lys	Val	Pro	Pro	Ile	Lys	Pro
65					70					75					80
Asn	Ala	Gly	Glu	Glu	Ser	Val	Met	Asn	Leu	Asp	Lys	Leu	Arg	Phe	Ala
				85					90					95	
Asp	Gly	Ser	Пe	Arg	Thr	Ser	Glu	Leu	Arg	Leu	Ser	Met	Gln	Lys	Ser
			100					105					110		
Met	Gln	Asn	His	Ala	Ala	Va]	Phe	Arg	Val	Gly	Ser	Val	Leu	Gln	Glu
		115					120					125			
Gly	Cys	Gly	Lys	He	Ser	Lys	Leu	Tyr	Gly	Asp	Leu	Lys	His	Leu	Lys
	130					135					140				
Thr	Phe	Asp	Arg	Gly	Met	Val	Trp	Asn	Thr	Asp	Leu	Val	Glu	Thr	Leu
145					150					155					160
Glu	Leu	Gln	Asn	Leu	Met	Leu	Cys	Ala	Leu	Gln	Thr	He	Tyr	Gly	Ala
				165					170					175	
Glu	Ala	Arg	Lys	Glu	Ser	Arg	Gly	Ala	His	Ala	Arg	Glu	Asp	Tyr	Lys
			180					185					190		
Va]	Arg	He	Asp	Glu	Tyr	Asp	Tyr	Ser	Lys	Pro	lle	Gln	Gly	Gln	Gln
		195					200					205			
Lys	Lys	Pro	Phe	Glu	Glu	His	Trp	Arg	Lys	His	Thr	Leu	Ser	Tyr	Val
	210					215					220				
Asp	Val	G1y	Thr	Gly	Lys	Val	Thr	Leu	Glu	Tyr	Arg	Pro	Val	He	Asp
225					230					235					240
Lys	Thr	Leu	Asn	Glu	Ala	Asp	Cys	Ala	Thr	Val	Pro	Pro	Ala	lle	Arg
				245					250					255	
Ser	Tyr														

<211> 149

<212> PRT

<213> Homo sapiens

<400> 576

Met Tyr Met Glu Pro Ser Ala Arg Ser Ala Gly Tyr Ser Pro Gln Gln

1 5 10 15

Lys Arg Leu Ser Lys Thr Arg 11e Pro Arg Leu Gln Ser Leu Thr Asp 20 25 30

Gln Ala Ala Leu Trp Gly Thr Thr Cys Asp Gln Val Asn Ala Lys Gln 35 40 45

Gly Pro Lys Pro Ser Pro Gly His Arg Leu Arg Arg Asn Leu Pro Gly 50 55 60

Glu Lys Trp Glu Ile Asp Phe Thr Lys Val Lys Pro His Gln Ala Gly
65 70 75 80

Tyr Lys Tyr Leu Leu Val Leu Val Asp Thr Phe Ser Gly Trp Thr Glu 85 90 95

Ala Phe Ala Thr Lys Asn Glu Thr Ala Asn Leu Val Val Lys Phe Leu
100 105 110

Leu Asn Glu Ile Ile Pro Arg Tyr Gly Leu Pro Ala Ala Ile Gly Ser 115 120 125

Asp Asn Gly Pro Ala Phe Thr Ser Ser Ile Val Leu Ser Val Ser Lys 130 135 140

Ala Leu Asn Ile Gln

145

<210> 4954

<211> 602

<212> PRT

<213> Homo sapiens

<400> 577

Met Glu Ser Gln Ala Thr Ser Ala Ser IIe Asn Asn Ser Asn Pro Ser

1 5 10 15

Thr Ser Glu Gln Ala Ser Asp Thr Ala Ser Ala Val Thr Ser Ser Gln 20 25 30

Pro Ser Thr Val Ser Glu Thr Ser Ala Thr Leu Thr Ser Asn Ser Thr

		35					40					45			
Thr	Gly	Thr	Ser	He	Gly	Asp	Asp	Ser	Arg	Arg	Thr	Thr	Ser	Ser	Ala
	50					55					60				
Val	Thr	Glu	Thr	Gly	Pro	Pro	Ala	Met	Pro	Arg	Leu	Pro	Ser	Cys	Cys
65					70					75					80
Pro	G1n	His	Ser	Pro	Cys	Gly	Gly	Ser	Ser	Gln	Asn	His	His	Ala	Leu
				85					90					95	
Gly	His	Pro	His	Thr	Ser	Cys	Phe	Gln	Gln	His	Gly	His	His	Phe	Gln
			100					105					110		
His	His	His	His	His	His	His	Thr	Pro	His	Pro	Ala	Val	Pro	Val	Ser
		115					120					125			
Pro	Ser	Phe	Ser	Asp	Pro	Λla	Cys	Pro	Val	Glu	Arg	Pro	Pro	Gln	Val
	130					135					140				•
	Ala	Pro	Cys	Gly	Ala	Asn	Ser	Ser	Ser	Gly	Thr	Ser	Tyr	His	Glu
145					150					155					160
Gln	Gln	Ala	Leu		Val	Asp	Leu	Ser		Ser	Gly	lle	Arg		His
				165					170					175	
Gly	Ser	Gly		Phe	His	Gly	Ala		Ala	Phe	Asp	Pro		Cys	Pro
			180					185	0.7		0.1		190		
Val	Ser		Ser	Arg	Ala	Ala		Phe	Gly	His	GIn		Ala	Ala	Ala
A 1	D	195	C1	D	1	C	200	11.	Λ	C1	т	205	C	C	M . 4
Ala		ser	GIN	Pro	Leu	Ser	ser	116	Asp	GIY		GIY	ser	ser	Met
Vol.	210	Cln	Dwo	C1n	Dro	215 Cln	Dno	Drag	Dro	Cln	220 Pro	Son	Lou	Son	San
Val 225	MIa	OIII	110	0111	230	Gln	110	110	110	235	110	261	Leu	Se1	240
	Arø	His	Tyr	Met		Pro	Pro	Tyr	Ala		Len	Thr	Arg	Pro	
Cyo	6	11.10	. , .	245			110	.,.	250		Lou		111 6	255	Bea
His	His	Gln	Ala			Cys	Pro	His			Glv	Asn	Pro		Pro
			260			-,		265					270		
Gln	Thr	Gln		Pro	Pro	Gln	Val		Tyr	Va]	lle	Pro		Pro	Val
		275					280	•	•			285			
His	Ala	Phe	His	Ser	Gln	11e	Ser	Ser	His	Ala	Thr	Ser	His	Pro	Val
	290					295					300				
Ala	Pro	Pro	Pro	Pro	Thr	His	Leu	Ala	Ser	Thr	Ala	Ala	Pro	lle	Pro
305					310					315					320
G1n	His	Leu	Pro	Pro	Thr	llis	Gln	Pro	He	Ser	His	His	He	Pro	Ala

				325					330					335	
Thr	Ala	Pro	Pro	Ala	Gln	Arg	Leu	His	Pro	His	Glu	Val	Met	Gln	Arg
			340					345					350		
Met	Glu	Va]	Gln	Arg	Arg	Arg	Met	Met	Gln	His	Pro	Thr	G]y	Leu	Phe
		355					360					365			
Val	Phe	Cys	Val	Ser	Arg	Arg	Ala	His	Glu	Arg	Pro	Pro	Pro	His	P.ro
	370					375					380				
His	Arg	Met	His	Pro	Asn	Tyr	Gly	His	Gly	His	His	lle	His	Val	Pro
385					390					395					400
Gln	Thr	Met	Ser	Ser	His	Pro	Arg	Gln	Ala	Pro	Glu	Arg	Ser	Ala	Trp
				405					410					415	
Glu	Leu	Gly	lle	Glu	Ala	Gly	Val	Thr	Ala	Ala	Thr	Tyr	Thr	Pro	Gly
			420					425					430		
Ala	Leu	His	Pro	His	Leu	Ala	His	Tyr	His	Ala	Pro	Pro	Arg	Leu	His
		435					440					445			
His	Leu	Gln	Leu	Gly	Ala	Leu	Pro	Leu	Met	Val	Pro	Asp	Met	Ala	Gly
	450					455					460				
Tyr	Pro	His	lle	Arg	Tyr	lle	Ser	Ser	Gly	Leu	Asp	Gly	Thr	Ser	Phe
465					470					475					480
Arg	Gly	Pro	Phe	Arg	Gly	Asn	Phe	Glu	Glu	Leu	lle	His	Leu	Glu	Glu
				485					490					495	
Arg	Leu	G] y	Asn	Val	Asn	Arg	Gly	Ala	Ser	Gln	G1 y	Thr	He	Glu	Arg
			500					505					510		
Cys	Thr	Tyr	Pro	His	Lys	Tyr	Lys	Lys	Val	Thr	Thr	Asp	Trp	Phe	Ser
		515					520					525			
Gln	Arg	Lys	Leu	His	Cys	Lys	Gln	Asp	Gly	Glu	Glu	Gly	Thr	Glu	Glu
	530					535					540				
Asp	Thr	Glu	Glu	Lys	Cys	Thr	lle	Cys	Leu	Ser	He	Leu	Glu	Glu	Gly
545					550					555					560
Glu	Asp	Val	Arg	Arg	Leu	Pro	Cys	Met	His	Leu	Phe	His	Gln	Val	Cys
				565					570					575	
Val	Asp	Gln		Leu	lle	Thr	Asn		Lys	Cys	Pro	He	Cys	Arg	Val
			580					585					590		
Asp	He		Ala	Gln	Leu	Pro	Ser	Glu	Ser						
		595					600								

<210> 4955 <211> 116 <212> PRT <213> Homo sapiens <400> 578 Met Val Ser Gly Met Gly Met Gly Asn Arg Asp Thr Asn Pro Asn Gly 1 10 15 Lys Pro Leu Ala Lys Gln Pro Pro Glu Phe Val Leu Ile Val Tyr Ser 25 Gln Ser Leu Ala Thr Ala His Thr Leu Phe Phe Ser Tyr Lys Gln Lys 45 Glu Leu Ser Leu Ser Ala Met Asn Pro Ala Ile Pro Arg Lys Lys Ala 50 55 Asn Ala Leu Ala Ser Ser Pro Val Arg Ala Thr His Ser Ile Ser Thr 70 75 Phe Cys Met Leu Lys Leu Cys His Arg Arg Arg Ala Ser Ala His Asp 85 90 Gln Phe Phe Phe Trp Ser lle Gly Ser Phe Cys Leu Arg Ile Phe Val 100 105 110 Cys Val Tyr Leu 115 <210> 4956 <211> 376 <212> PRT <213> Homo sapiens <400> 579 Met Ala Asp Ser Arg Arg Val 11e 11e Ala Ser Trp Tyr Arg Thr Phe 10

Met Gly Ile Val Asn Leu Phe Gly Leu Glu Thr Lys Thr Cys Trp Asn

Val Thr Arg Ile Glu Pro Leu Asn Glu Val Gln Ser Cys Glu Gly Leu

30

		35					40					45			
Arg	Asp	Pro	Ala	Cys	Phe	Tyr	Val	Gly	Val	lle	Phe	Ile	Leu	Asn	Gly
	50					55					60				
Leu	Met	Met	Gly	Leu	Phe	Phe	lle	Tyr	Gly	Thr	Tyr	Leu	Ser	Gly	Thr
65					70					75					80
G] u	Leu	Gly	Gly	Leu	He	Thr	Val	Leu	Cys	Phe	Phe	Phe	Asn	His	Gly
				85					90					95	
Glu	Ala	Thr	Cys	Val	Met	Trp	Thr	Pro	Pro	Leu	Arg	Glu	Ser	Phe	Ser
			100					105					110		
Tyr	Pro	Phe	Leu	Val	Leu	Gln	Met	Tyr	Val	Leu	Thr	Leu	Ile	Leu	Arg
		115					120					125			
Thr	Ser	Ser	Asn	Asp	Arg	Arg	Pro	Phe	He	Ala	Leu	Cys	Leu	Ser	Asn
	130					135					140				
	Ala	Phe	Met	Leu	Pro	Trp	Gln	Phe	A]a	G] n	Phe	He	Leu	Phe	Thr
145					150					155					160
Gln	He	Ala	Ser		Phe	Pro	Met	Tyr		Va]	Gly	Tyr	He		Pro
_				165			_	.0.	170			_		175	
Ser	Lys	Phe		Lys	He	He	Tyr		Asn	Met	He	Ser		Thr	Leu
C	ĎΙ	7.1	180		DI	6.1		185		Tr.		0	190	T	an.
Ser	Phe	He	Leu	Met	Phe	Gly		Ser	Met	lyr	Leu		Ser	Tyr	Tyr
C	C	195	1	1	M-4	тъ	200 T	A 1 -	т1.	33.	1	205			C1
ser		Ser	Leu	Leu	Met		irp	Ala	.11e	11e		Lys	Arg	Asn	Glu
Па	210	Lys	Lou	Clv	Val	215 Sor	Lve	Lou	Acn	Cve	220 Trp	Lou	110	Cln	C1v
225	OIII	Lys	Leu	O1 y	230	261	Lys	Leu	изп	235	пр	Leu	116	0111	240
	Ala	Trp	Trn	Cvs		Thr	He	116	Len		Phe	Leu	Thr	Ser	
001	7110	11 15	пр	245	013	1111	110	110	250		THE	Lea	1111	255	Lys
He	Leu	Gly	Val	Ser	Asp	His	11e	Cvs			Asp	Leu	He		Ala
			260		,			265				,	270		
Gly	Thr	Leu	Arg	Tyr	Thr	Asp	Phē		Thr	Leu	Lvs	Tyr		Cys	Ser
		275		·			280	•			•	285		•	
Pro	Glu	Phe	Asp	Phe	Met	Glu	Lys	Ala	Thr	Leu	Leu		Tyr	Thr	Lys
	290					295					300				
Thr	Leu	Leu	Leu	Pro	Val	Va1	Met	Val	Пе	Thr	Cys	Phe	He	Phe	Lys
305					310					315					320
Lvs	Thr	Val	G] v	Asp	He	Ser	Arg	Val	l.eu	Ala	Thr	Asn	Val	Tvr	Leu

| Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | Sample | S

<400> 580

<213> Homo sapiens

Met Tyr Pro Ala Leu Leu Val Pro Ser Ser Ser Pro Ser Ser Val Phe

1 5 10 15

Gly Val Pro Glu Thr Gly Phe Ser Gln Trp Pro Gly Pro Gln Gly Asn
20 25 30

Cys Arg Val 11e Gln Ser Gly Ser Ala Ser Leu Gln Gly Pro Leu Pro 35 40 45

Gly Arg Ala Ser Trp His Pro Ala 11e Gly Gly Pro Asp Thr Pro Ala 50 55 60

Arg Asp Pro Ala Thr Gln Glu Val Pro Thr Pro Ser Gln Ser Asp Pro 65 70 75 80

Gly Pro Arg Ala Asp Arg Ser Pro Pro Leu Lys Arg Leu Leu Pro Leu 85 90 95

Tyr Pro Thr Pro Arg Asp Ser Phe His Ser Gly 11e Ser 100 105

<210> 4958

<211> 363

<212> PRT

<213> Homo sapiens

<400)> 58	31													
Met	Pro	Val	Arg	Thr	Leu	Phe	Pro	Glu	Ser	Trp	Leu	Trp	Arg	Lys	Phe
1				5					10					15	
Thr	Leu	Pro	Lys	Ser	Lys	Ser	Gly	He	Ser	His	Tyr	Pro	Пе	Ser	Val
			20					25					30		
Lys	Val	Pro	Лѕр	Ser	He	Thr	Thr	Trp	Gln	Phe	Val	Val	Val	Ser	Leu
		35					40					45			
Lys	Ala	Gly	Gln	Gly	Leu	Cys	Val	Ser	Asp	Pro	Phe	Glu	Leu	Thr	Val
	50					55					60				
Met	Lys	Ser	Phe	Phe	Val	Asp	Leu	Lys	Leu	Pro	Ser	Ser	Val	lle	Arg
65					70					75					80
Asn	Glu	Gln	Val	Gln	lle	Gln	Ala	Met	Leu	Tyr	Asn	Phe	Arg	Asp	Arg
				85					90					95	
Gln	Ala	Lys	Val	Arg	Val	Glu	Phe	Pro	His	Lys	Glu	Thr	Leu	Cys	Ser
			100					105					110		
Ala	Ser	Lys	Pro	Gly	Ala	Pro	Ser	His	Gln	Ala	Gly	Val	Gln	Ile	Gln
		115					120					125			
Gln	Thr	Ser	Tyr	Ser	Ile	Val	Leu	Glu	Pro	Gln	Gly	Gln	Thr	Gln	Thr
	130					135					140				
Lys	Leu	Val	Pro	Arg	Gln	Glu	Phe	Leu	Asn	Met	Val	Pro	Asp	Thr	Glu
145					150					155					160
Ala	Glu	Val	Phe	He	Ser	Val	Gln	Gly	Tyr	Thr	Gln	Met	Leu	Thr	His
				165					170					175	
Arg	Ser	Ser		Gly	Thr	Tyr	His	Thr	Ser	Lys	Gly	Asn	Pro	Gly	Ser
			180					185					190		
Thr	Trp	Leu	Thr	Ser	Tyr	Val		Arg	Val	Phe	Ala	Leu	Ala	Tyr	Ser
		195					200					205			
Met		Thr	Thr	Gln	Val		Ser	Leu	Ser	Ser		Cys	Asp	Met	Ala
	210					215					220				
Asn	Trp	He	He	He		Arg	GIn	Ala	Glu	Asp	Gly	His	Phe	Leu	Glu
225					230					235					240
Lys	G] y	Pro	Val	Val	Met	Thr	Ser	Met	Gln	Asp	Gly	Tyr	Gln	G] y	Ser
				245					250					255	
Glu	Glu	Asp	Val	Ser	Leu	Thr	Ala	Leu	Val	Leu	He	Ala	Leu	Λsn	Glu
			260					265					270		

Gly Lys Glu Leu Cys Arg Gln Lys Asn Leu Met Ala Ser Ile Glu Arg Ala Arg Gly Phe Leu Glu Arg Lys Leu Pro Asp Ile Gln Thr Thr Phe Ala Val Ala Ile Ala Ser Tyr Ala Leu Ala Leu Ala Asn Ser Ser GIn Ala Asn Asp Cys Leu Asp Ser Phe Ala Ser Pro Ser Gly Cys Gly Met Leu Leu Asn Gln Pro Gln Ser Trp Ser Gly Glu Gly Val Ile Ser Asn Pro Ala Met Cys Tyr Ser Ser Leu Ser Val Ser

<210> 4959

<211> 359

<212> PRT

<213> Homo sapiens

<400> 582

Met Pro Pro Thr Ser Ser Thr Pro Ser Leu Ser Arg Pro Gly Leu Gly Gln Ala Gly Lys Pro Asp Thr Gly Ser His Pro Pro Pro Thr Ile Ser Thr Ser Ile Phe Leu Ser Cys Phe Pro Thr Ile Pro Leu Ser Arg Pro Arg Thr Thr Gly Pro Ser His Ser Tyr Gln Ser He Ser His Pro Arg Ser Cys Arg Asp Val Pro Asp Asp Ile Gln Pro 11e Thr Ser Leu Pro Gly Val Ala Arg Tyr Gly Val Lys Arg Leu Glu Glu Met Leu Arg Pro Leu Val Glu Glu Gly Leu Arg Cys Val Leu Ile Phe Gly Val Pro Ser Arg Val Pro Lys Asp Glu Arg Gly Ser Ala Ala Asp Ser Glu Glu Ser

Pro	Ala	Ile	Glu	Ala	Ile	His	Leu	Leu	Arg	Lys	Thr	Phe	Pro	Asn	Leu
	130					135					140				
Leu	Val	Ala	Cys	Asp	Val	Cys	Leu	Cys	Pro	Tyr	Thr	Ser	His	Gly	His
145					150					155					160
Cys	Gly	Leu	Leu	Ser	Glu	Asn	Gly	Ala	Phe	Arg	Ala	Glu	Glu	Ser	Arg
				165					170					175	
Gln	Arg	Leu	Ala	Glu	Val	Ala	Leu	Ala	Tyr	Ala	Lys	Ala	Gly	Cys	Gln
			180					185					190	,	
Val	Val	Ala	Pro	Ser	Asp	Met	Met	Asp	Gly	Arg	Val	Glu	Ala	lle	Lys
		195					200					205			
Glu	Ala	Leu	Met	Ala	His	Gly	Leu	Gly	Asn	Arg	Val	Ser	Val	Met	Ser
	210					215					220				
Tyr	Ser	Ala	Lys	Phe	Ala	Ser	Cys	Phe	Tyr	Gly	Pro	Phe	Arg	Asp	Ala
225					230					235					240
Ala	Lys	Ser	Ser	Pro	Ala	Phe	Gly	Asp	Arg	Arg	Cys	Tyr	Gln	Leu	Pro
				245					250					255	
Pro	Gly	Ala	Arg	Gly	Leu	Ala	Leu	Arg	Ala	Val	Asp	Arg	Asp	Val	Arg
			260					265					270		
Glu	Gly	Ala	Asp	Met	Leu	Met	Val	Lys	Pro	Gly	Met	Leu	Tyr	Leu	Asp
		275					280					285			
He	Val	Arg	Glu	Va]	Lys	Asp	Lys	His	Pro	Asp	Leu	Pro	Leu	Ala	Val
	290					295			`		300				
Tyr	His	Val	Ser	Gly	Glu	Phe	Ala	Met	Leu	Trp	His	Gly	Ala	Gln	Ala
305					310					315					320
Gly	Ala	Phe	Asp	Leu	Lys	Ala	Ala	Val	Leu	Glu	Ala	Met	Thr	Λla	Phe
				325					330					335	
Arg	Arg	Ala	Gly	Ala	Asp	He	lle	lle	Thr	Tyr	Tyr	Thr	Pro	Gln	Leu
			340					345					350		
Leu	Gln	Trp	Leu	Lys	Glu	Glu									
		355													

<211> 508

<212> PRT

<213> Homo sapiens

<400)> 58	33													
Met	Met	Trp	Gly	Gly	Gly	Leu	Asp	Leu	Cys	Pro	Met	Pro	Gly	Gln	Leu
i				5					10					15	
Lys	Phe	Pro	Pro	Cys	Leu	Ser	Arg	Cys	Leu	Leu	Trp	Glu	Pro	Pro	Ser
			20					25					30		
Leu	Tyr	Leu	Thr	Gln	Pro	Thr	Ser	Ser	Leu	Ala	Glu	Pro	Gln	Ala	Leu
		35					40					45			
Ile	Cys	Met	Thr	Ser	Ser	Ser	Ser	Gly	Leu	Phe	Ile	Gln	Asp	Asp	Asn
	50					55					60				
Met	Glu	Lys	Leu	Glu	Glu	Ile	Ile	Glu	Lys	Tyr	Pro	Arg	Ala	Phe	Pro
65					70					75					80
Phe	Trp	11e	Gly	Pro	Phe	Gln	Ala	Phe	Phe	Cys	lle	Tyr	Asp	Pro	Asp
				85					90					95	
Tyr	Ala	Lys	Thr	Leu	Leu	Ser	Arg	Thr	Asp	Pro	Lys	Ser	Gln	Tyr	Leu
			100					105					110		
Gln	Lys	Phe	Ser	Pro	Pro	Leu	Leu	Gly	Lys	Gly	Leu	Ala	Ala	Leu	Asp
		115					120					125			
Gly	Pro	Lys	Trp	Phe	Gln	His	Arg	Arg	Leu	Leu	Thr	Pro	G1 y	Phe	His
	130					135					140				
Phe	Asn	Ile	Leu	Lys	Ala	Tyr	He	Glu	Val	Met	Ala	His	Ser	Val	Lys
145					150					155					160
Met	Met	Leu	Asp		Trp	Glu	Lys	He		Ser	Thr	Gln	Asp		Ser
				165					170					175	
Val	Glu	Val		Glu	His	lle	Asn		Met	Ser	Leu	Asp	He	Val	Met
			180					185					190		
Lys	Cys		Phe	Ser	Lys	Glu		Asn	Cys	Gln	Thr		Ser	Thr	His
	_	195					200			_	_	205			
Asp		Tyr	Ala	Lys	Ala		Phe	Glu	Leu	Ser		He	He	Phe	His
	210	_				215		0		~ 1	220	631			
	Leu	Tyr	Ser	Leu		Tyr	His	Ser	Asp		He	Phe	Lys	Leu	
225	61	C.1	т		230	C1			C	235	12 1			C1	240 T
Pro	GIn	Gly	lyr		Phe	GIn	Lys	Leu		Arg	Val	Leu	Asn		lyr
TI		TI		245	C.I	C 1		,	250	C		C I	ΛĪ.	255	W 1
Ihr	Asp	Ihr		116	GIn	ьIu	Arg		Lys	5er	Leu	6In	Ala	61 y	val
			260					265					270		

Lys	Gln	Asp	Asn	Thr	Pro	Lys	Arg	Lys	Tyr	Gln	Asp	Phe	Leu	Asp	He
		275					280	•				285			
Val	Leu	Ser	Λla	Lys	Asp	Glu	Ser	G1 y	Ser	Ser	Phe	Ser	Asp	He	Asp
	290					295					300				
Val	His	Ser	Glu	Val	Ser	Thr	Phe	Leu	Leu	Ala	G]y	His	Asp	Thr	Leu
305					310					315					320
Ala	Ala	Ser	He	Ser	Trp	11e	Leu	Týr	Cys	Leu	Ala	Leu	Asn	Pro	Glu
				325					330					335	
His	Gln	Glu	Arg	Cys	Arg	Glu	Glu	Val	Arg	Gly	Ile	Leu	Gly	Asp	Gly
			340					345					350		
Ser	Ser	Ile	Thr	Trp	Asp	Gln	Leu	Gly	Glu	Met	Ser	Tyr	Thr	Thr	Met
		355					360					365			
Cys	Пе	Lys	Glu	Thr	Cys	Arg	Leu	Πe	Pro	Ala	Val	Pro	Ser	lle	Ser
	370					375					380				
Arg	Asp	Leu	Ser	Lys	Pro	Leu	Thr	Phe	Pro	Asp	Gly	Cys	Thr	Leu	Pro
385					390					395					400
Ala	Gly	Ile	Thr	Val	Val	Leu	Ser	Ile	Trp	Gly	Leu	His	His	Asn	Pro
				405					410					415	
Ala	Val	Trp	Lys	Asn	Pro	Lys	Val	Phe	Asp	Pro	Leu	Arg	Phe	Ser	Gln
			420					425					430		
Glu	Asn	Ser	Asp	Gln	Arg	His	Pro	Tyr	Ala	Tyr	Leu	Pro	Phe	Ser	Ala
		435					440					445			
Gly	Ser	Arg	Asn	Cys	He	Gly	Gln	Glu	Phe	Ala	Met	He	G] u	Leu	Lys
	450					455					460				
Val	Thr	He	Ala	Leu	Ile	Leu	Leu	His	Phe	Arg	Val	Thr	Pro	Asp	Pro
465					470					475					480
Thr	Arg	Pro	Leu	Thr	Phe	Pro	Asn	His	Phe	Ile	Leu	Lys	Pro	Lys	Asn
				485					490					495	
Gly	Met	Tyr	Leu	His	Leu	Lys	Lys	Leu	Ser	Glu	Cys				
			500					505							

<211> 1264

<212> PRT

<213> Homo sapiens

<40	0> 5	84													
Met	Cys	Lys	Lys	Leu	Thr	Lys	Leu	Ala	Lys	Glu	Asn	Asp	Ser	Met	Lys
1				5					10					15	
Glu	Glu	Leu	Leu	Lys	Tyr	Arg	Ser	Leu	Tyr	Gly	Asp	Leu	Asp	Ser	Ala
			20					25					30		
Leu	Ser	Ala	Glu	Glu	Leu	Ala	Asp	Ala	Pro	His	Ser	Arg	Glu	Thr	Glu
		35					40					45			
Leu	Lys	Val	His	Leu	Lys	Leu	Val	Glu	Glu	Glu	Ala	Asn	Leu	Leu	Ser
	50					55					60				
Arg	Arg	Ile	Val	Glu	Leu	Glu	Val	Glu	Asn	Arg	Gly	Leu	Arg	Ala	Glu
65					70					75					80
Met	Asp	Asp	Met	Lys	Asp	His	Gly	Gly	Gly	Cys	Gly	Gly	Pro	Glu	Ala
				85					90					95	
Arg	Leu	Ala	Phe	Ser	Ala	Leu	Gly	G1 y	Gly	Glu	Cys	Gly	Glu	Ser	Leu
			100					105					110		
Ala	Glu	Leu	Arg	Arg	His	Leu	Gln	Phe	Val	Glu	Glu	Glu	Ala	Glu	Leu
		115					120					125			
Leu		Arg	Ser	Ser	Ala	Glu	Leu	G] u	Asp	Gln	Asn	Lys	Leu	Leu	Leu
	130					135					140				
	Glu	Leu	Ala	Lys		Arg	Ser	Glu	His		Leu	Asp	Val	Ala	Leu
145	0.1				150			_		155	_				160
Ser	Glu	Asp	Ser		Ser	Val	Leu	Ser		Pro	Ser	Gln	Glu		Leu
4.1	4.7	4.1		165	C.1		C 1	0.1	170	0	0.1	,		175	
Ala	Ala	Ala		Leu	GIN	He	Gly		Leu	Ser	Gly	Lys		Lys	Lys
l	C1	т	180	Λ	Δ	V - 1	1	185	C	4	,	61	190	C	
Leu	OHI		Gju	ASII	АГВ	Val		Leu	ser	ASN	Leu		Arg	Cys	Asp
Lau	Λla	195 Sor	Cvc	Cln.	Sor	The	200	Dro	Mot	ارم ا	C1	205	1.00	۸1	C1
Leu	210	361	Cys	OIII	361	Thr 215	Alg	Lio	Met	Leu		1111	ASP	мта	GIU
Ala		Aen	Sor	Δla	Gln	Cys	Val	Pro	Λla	Pro	220	Cly	C1v	The	ui o
225	Oly	пар	261	Mia	230	Cys	vai	110	Ма	235	Leu	оту	O I U	1111	240
	Ser	His	Ala	Val		Leu	Cvs	Arg	Ala		Glu	Ala	Glu	Val	
				245	6	204	J, 5	8	250	8	J. U	. 1 L C	Jju	255	1,0 U
Pro	Glv	Leu	Arg		Gln	Ala	Ala	Leu		Ser	Lvs	Ala	He		Val
	,		260			-		265			-,-		270		

Leu	Val	A1a 275	Asp	Ala	Asn	Gly	280	Thr	Ala	Gly	Leu	Arg 285	Leu	Cys	Leu
Asn	Aen		Cvs	Ala	Aen	Pho	Arg	الم ا	Hic	Glu	Δla		Acn	Acn	Sor
пэр	Mon	Oru	Cy3	MIG	изр	THE	мв	Leu	1115	Oiu	Mid	110	nsp	nsn	361
	290					295					300				
Glu	Gly	Pro	Arg	Asp	Thr		Leu	lle	His	Ala		Leu	Val	Arg	Leu
305					310					315				J	320
Ser	Val	Leu	Gln	Gln	Glu	Leu	Asn	Ala	Phe	Thr	Arg	Lys	Ala	Asp	Ala
				325					330					335	
Val	Leu	Gly	Cys	Ser	Val	Lys	Glu	Gln	Gln	Glu	Ser	Phe	Ser	Ser	Leu
			340					345					350		
Pro	Pro	Leu	Gly	Ser	Gln	Gly	Leu	Ser	Lys	Glu	lle	Leu	Leu	Ala	Lys
		355					360					365			
Asp	Leu	Gly	Ser	Asp	Phe	Gln	Pro	Pro	Asp	Phe	Arg	Asp	Leu	Pro	Glu
	370					375					380				
Trp	Glu	Pro	Arg	lle	Arg	Glu	Ala	Phe	Arg	Thr	Gly	Asp	Leu	Asp	Ser
385					390					395					400
Lys	Pro	Asp	Pro	Ser	Arg	Ser	Phe	Arg	Pro	Tyr	Arg	Ala	Glu	Asp	Asn
				405					410					415	
Asp	Ser	Tyr	Ala	Ser	Glu	Ile	Lys	Glu	Leu	Gln	Leu	Val	Leu	Ala	Glu
			420					425					430		
Ala	His	Asp	Ser	Leu	Arg	Gly	Leu	Gln	Glu	Gln	Leu	Ser	Gln	Glu	Arg
		435					440					445			
Gln	Leu	Arg	Lys	Glu	Glu	Ala	Asp	Asn	Phe	Asn	Gln	Lys	Met	Val	Gln
	450					455					460				
Leu	Lys	Glu	Asp	Gln	Gln	Arg	Ala	Leu	Leu	Arg	Arg	Glu	Phe	Glu	Leu
465					470					475					480
Gln,	Ser	Leu	Ser		GIn	Arg	Arg	Leu		Gln	Lys	Phe	Trp		Gln
				485					490					495	
Glu	Lys	Asn		Leu	Val	GIn	Glu		Gln	Gln	Phe	Lys		Asn	Phe
			500					505	D)	,			510		
Leu	Leu		Phe	Met	Lys	Leu	Arg	lrp	Phe	Leu	Lys		Trp	Arg	GIn
C1	1	515	,	15	C	61	520			Di		525	** 1		0
огу		val	Leu	rro	ser		Gly	ASP	Asp	rne		61u	val	Asn	5er
Mo+	530	V - 1	Lou	Т.,,,,	Lau	535	Met	C1	C1	C1	540	11.	Λ	Д1.	C1
DIT: I	1.V.	val	1.150	1 V L	1 (-) [1	1 (3)	WH-3	13111	1 1 1 1 1	1 2 1 1 1	(, 1	110	ASD	1112	LIP

545	;				550)				555	5				560	
His	Ser	· Asp	Asr	Lys	Ala	Cys	Thr	Gly	/ Asp	Se ₁	· Trp	Thr	- Glr	n Asr	Thr	
				565					570					575		
Pro	Asr	ı Glu	туз	· Ile	Lys	Thr	Leu	Λlε	Asp	Met	Lys	. Val	Thr	Leu	Lys	
			580					585					590			
Glu	Leu	Cys	Trp	Leu	Leu	Arg	Asp	Glu	ιArg	g Arg	Gly	Leu	Thr	Glu	Leu	
		595	5				600					605	;			
Gln	Gln	Gln	Phe	Ala	Lys	Ala	Lys	Ala	Thr	Trp	Glu	Thr	Glu	Arg	Ala	
	610					615					620					
Glu	Leu	Lys	Gly	His	Thr	Ser	Gln	Met	Glu	Leu	Lys	Thr	Gly	Lys	Gly	
625					630					635					640	
Ala	Gly	Glu	Arg	Ala	Gly	Pro	Asp	Trp	Lys	Ala	Ala	Leu	Gln	Arg	Glu	
				645					650					655		
Arg	Glu	Glu	Gln	Gln	His	Leu	Leu	Ala	Glu	Ser	Tyr	Ser	Ala	Val	Met	
			660					665					670			
Glu	Leu	Thr	Arg	Gln	Leu	Gln	He	Ser	Glu	Arg	Asn	Trp	Ser	Gln	Glu	
		675					680					685				
Lys	Leu	Gln	Leu	Val	Glu	Arg	Leu	Gln	Gly	Glu	Lys	Gln	Gln	Val	Glu	
_	690					695					700					
Gln	Gln	Val	Lys	Glu	Leu	Gln	Asn	Arg	Leu	Ser	Gln	Leu	Gln	Lys	Ala	
705					710					715					720	
Ala	Asp	Pro	Trp	Va]	Leu	Lys	His	Ser	Glu	Leu	Glu	Lys	Gln	Asp	Asn	
				725					730					735		
Ser	Trp	Lys	Glu	Thr	Arg	Ser	Glu	Lys	He	His	Asp	Lys	Glu	Ala	Val	
			740					745					750			
Ser	Glu	Va]	Glu	Leu	Gly	G]y	Asn	Gly	Leu	Lys	Arg	Thr	Lys	Ser	Val	
		755					760					765				
Ser		Met	Ser	Glu	Phe	Glu	Ser	Leu	Leu	Asp	Cys	Ser	Pro	Tyr	Leu	
	770					775					780					
	Gly	Gly	Asp	Ala		Gly	Lys	Lys	Leu	Pro	Asn	Asn	Pro	Ala	Phe	
785					790					795					800	
Gly	Phe	Val	Ser	Ser	Glu	Pro	Gly	Asp	Pro	Glu	Lys	Asp	Thr	Lys	Glu	
				805					810					815		
_ys	Pro	Gly		Ser	Ser	Arg	Asp	Cys	Asn	His	Leu	Gly	Ala	Leu	Ala	
-			820					825					830			
.ys	GIn	Asp	Pro	Pro	Glv	Arg	G1n	Met	Gla	Aro	Ser	Tyr	Thr	Ala	Pro	

		835					840					845			
Asp	Lys	Thr	Gly	lle	Arg	Val	Tyr	Tyr	Ser	Pro	Pro	Val	Ala	Arg	Arg
	850					855					860				
Leu	Gly	Val	Pro	Val	Va]	His	Asp	Lys	Glu	Gly	Lys	He	lle	11e	Glu
865					870					875					880
Pro	Gly	Phe	Leu	Phe	Thr	Thr	Ala	Lys	Pro	Lys	Glu	Ser	Ala	Glu	Ala
				885					890					895	
Asp	Gly	Leu	Ala	Glu	Ser	Ser	Tyr	Gly	Arg	Trp	Leu	Cys	Asn	Phe	Ser
			900					965					910		
Arg	Gln	Arg	Leu	Asp	Gly	Gly	Ser	Ala	Gly	Ser	Pro	Ser	Ala	Ala	Gly
		915					920					925			
Pro	Gly	Phe	Pro	Ala	Ala	Leu	His	Asp	Phe	Glu	Met	Ser	Gly	Asn	Met
	930					935					940				
Ser	Asp	Asp	Met	Lys	Glu	He	Thr	Asn	Cys	Val	Arg	Gln	Ala	Met	Arg
945					950					955					960
Ser	Gly	Ser	Leu	Glu	Arg	Lys	Val	Lys	Ser	Thr	Ser	Ser	Gln	Thr	Val
				965					970					975	
G1 y	Leu	Ala	Ser	Val	Gly	Thr	Gln	Thr	Ile	Arg	Thr	Val	Ser	Val	Gly
			980					985					990		
Leu	Gln	Thr	Asp	Pro	Pro	Arg	Ser	Ser	Leu	His	Gly	Lys	Ala	Trp	Ser
		995					1000					005			
Pro	Arg	Ser	Ser	Ser	Leu	Val	Ser	Va]	Arg	Ser	Lys	Gln	He	Ser	Ser
	1010]	1015					1020				
Ser	Leu	Asp	Lys	Val	His	Ser	Arg	lle	Glu	Arg	Pro	Cys	Cys	Ser	Pro
1025	5				030					1035]	1040
Lys	Tyr	Gly	Ser	Pro	Lys	Leu	Gln	Arg	Arg	Ser	Val	Ser	Lys	Leu	Asp
]	1045]	1050					1055	
Ser	Ser	Lys	Asp	Arg	Ser	Leu	Trp	Asn	Leu	His	Gln	Gly	Lys	Gln	Asn
			1060				3	1065					1070		
Gly	Ser	Ala	Trp	Ala	Arg	Ser	Thr	Thr	Thr	Arg	Asp	Ser	Pro	Val	Leu
		1075					080					1085			
Arg	Asn	He	Asn	Asp	Gly	Leu	Ser	Ser	Leu	Phe	Ser	Val	Val	Glu	His
	090				1	1095					1100				
Ser	Gly	Ser	Thr	Glu	Ser	Val	Trp	Lys	Leu	G1 y	Met	Ser	Glu	Thr	Arg
1105	5				1110					1115]	1120
Ala	lve	Pro	Glu	Pro	Pro	lvs	Tyr	G1v	He	Va1	Gln	Glu	Phe	Phe	Arg

Asn Ala Cys Gly Arg Ala Pro Ser Pro Thr Ser Ser Ala Gly Glu Glu Gly Thr Lys Lys Pro Glu Pro Leu Ser Pro Ala Ser Tyr His Gln Pro Glu Gly Val Ala Arg lle Leu Asn Lys Lys Ala Ala Lys Leu Gly Ser Ser Glu Glu Val Arg Leu Thr Met Leu Pro Gln Val Gly Lys Asp Gly Val Leu Arg Asp Gly Asp Gly Ala Val Leu Pro Asn Glu Asp Ala Val Cys Asp Cys Ser Thr Gln Ser Leu Thr Ser Cys Phe Ala Arg Ser Ser Arg Ser Ala Ile Arg His Ser Pro Ser Lys Cys Arg Leu His Pro Ser Glu Ser Ser Trp Gly Gly Glu Glu Arg Ala Leu Pro Pro Ser Glu

<210> 4962

<211> 633

<212> PRT

<213> Homo sapiens

<400> 585

Met Glu Ser Gly Pro Lys Met Leu Ala Pro Val Cys Leu Val Glu Asn Asn Asn Glu Gln Leu Leu Val Asn Gln Gln Ala lle Gln lle Leu Glu Lys Ile Ser Gln Pro Val Val Val Ala Ile Val Gly Leu Tyr Arg Thr Gly Lys Ser Tyr Leu Met Asn His Leu Ala Gly Gln Asn His Gly Phe Pro Leu Gly Ser Thr Val Gln Ser Glu Thr Lys Gly 11e Trp Met

Trp	Cys	Val	Pro	His	Pro	Ser	Lys	Pro	Asn	His	Thr	Leu	Val	Leu	Leu
				85					90					95	
Asp	Thr	Glu	Gly	Leu	Gly	Asp	Val	Glu	Lys	Gly	Asp	Pro	Lys	Asn	Asp
			100					105					110		
Ser	Trp	11e	Phe	Ala	Leu	Ala	Val	Leu	Leu	Cys	Ser	Thr	Phe	Val	Tyr
		115					120					125			
Asn	Ser	Met	Ser	Thr	lle	Asn	His	Gln	Ala	Leu	Glu	Gln	Leu	His	Tyr
	130					135					140				
Val	Thr	Glu	Leu	Thr	Glu	Leu	Ile	Lys	Ala	Lys	Ser	Ser	Pro	Arg	Pro
145					150					155					160
Asp	Gly	Val	Glu	Asp	Ser	Thr	Glu	Phe	Val	Ser	Phe	Phe	Pro	Asp	Phe
				165					170					175	
Leu	Trp	Thr	Val	Arg	Asp	Phe	Thr	Leu	Glu	Leu	Lys	Leu	Asn	Gly	His
			180					185					190		
Pro	He	Thr	Glu	Asp	Glu	Tyr	Leu	Glu	Asn	Ala	Leu	Lys	Leu	He	G1n
		195					200					205			
Gly	Asn	Asn	Pro	Arg	Val	Gln	Thr	Ser	Asn	Phe	Pro	Arg	Glu	Cys	lle
	210					215					220				
Arg	Arg	Phe	Phe	Pro	Lys	Arg	Lys	Cys	Phe	Val	Phe	Asp	Arg	Pro	Thr
225					230					235					240
Asn	Asp	Lys	Asp		Leu	Ala	Asn	lle	Glu	Lys	Val	Ser	Glu	Lys	GIn
				245			•		250					255	
Leu	Asp	Pro		Phe	G1n	Glu	Gln		Asn	lle	Phe	Cys	Ser	Tyr	He
			260					265					270		
Phe	Thr		Ala	Arg	Thr	Lys		Leu	Arg	Glu	Gly		Thr	Val	Thr
0.3		275	_				280					285			
GTy		Arg	Leu	Gly	Thr	Leu	Ala	Val	Thr	Tyr		Glu	Ala	lle	Asn
c	290			Б	0	295	0.1				300				
	61y	Ala	Val	Pro		Leu	Glu	Asn	Ala		lle	Thr	Leu	Ala	
305	C1		C		310		G1			315		_	_	_	320
Arg	GIU	Asn	Ser		Ala	Val	GIn	Arg		Ala	Asp	Tyr	Tyr		Gln
C L n	Mad	41 a	C1	325	V - 1	1		n	330		T)	,	0.1	335	
oin	wet	ита		лrg	val	Lys	reu		ınr	Asp	Ihr	Leu		ыlы	Leu
Lou	A 015	Mat	340	A1	۸ l -	C	C1	345	C1	A 1	7.1	۸.1	350	DI	
ren	иsb		ms	ита	ита	Cys		Arg	01 u	АТА	116		11e	rne	Met
		355					360					365			

Glu	His 370	Ser	Phe	Lys	Asp	G1u 375	Asn	GIn	Glu	Phe	G1n 380	Lys	Lys	Phe	Met
Glu	Thr	Thr	Met	Asn	lve		Glv	Aen	Pho	Lou		Gla	Acn	Glu	61u
385	1111	111.2	inc c	71511	390	Lys	Oly	пар	1110	395	Leu	OIH	Non	oru	400
	Ser	Val	Gln	Tvr		Gln	Ala	lve	الم ا		Glu	Lau	Sor	Lve	
JCI	00,	101	OIII	405	Cys	OIII	MIG	Lys	410	11511	oru	Leu	261	415	Ory
Lou	Met	Glu	Sor		Sor	Λla	C1v	Sor		Sor	Val	Pro	C1v		Hic
LCu	MC C	010	420	116	261	MIG	01 y	425	1116	561	vai	110	430	ОГУ	1112
Lve	Leu	Tur		Clu	Thr	Lvc	Clu		Πla	Clu	Cln	Aan		Twn	Cln
Lys	Leu	435	MCC	Olu	1111	Lys	440	ni g	116	Olu	OIII	445	1 9 1	пр	GIII
Val	Pro		Lve	Glv	Val	lve		lve	Glu	Val	Pho		Ara	Pho	Lou
141	450	мб	Lys	Oly	141	455	MIG	Lys	Olu	vai	460	OIII	лід	THE	Leu
Glu	Ser	Gln	Met	Val	ء ا ا		Glu	Sor	110	ا ما		Sar	Acn	lve	Λla
465	501	OIII	,nc c	,01	470	0.14	Old	501	110	475	0111	501	пэр	Lys	480
	Thr	Asp	Arg	Glu		Ala	Val	Ala	Val		Arø	Ala	lvs	lvs	
200			6	485	D, O			1110	490	пор	6	ma	Lyo	495	014
Ala	Ala	Glu	Lvs		Gln	Glu	Leu	Leu		Gln	Lvs	Leu	Gln		Gln
			500					505			_, -		510		
Gln	Gln	Gln	Met	Glu	Ala	Gln	Asp		Ser	Arg	Lvs	Glu		He	Ala
		515					520	•			•	525			
Gln	Leu	Lys	Glu	Lys	Leu	Gln	Met	Glu	Arg	Glu	His	Leu	Leu	Arg	Glu
	530					535					540				
Gln	11e	Met	Met	Leu	G]u	His	Thr	Gln	Lys	Val	Gln	Asn	Asp	Trp	Leu
545					550					555					560
His	Glu	Gly	Phe	Lys	Lys	Lys	Tyr	Glu	Glu	Met	Asn	Ala	Glu	Пe	Ser
				565					570					575	
Gln	Phe	Lys	Arg	Met	He	Asp	Thr	Thr	Lys	Asn	Asp	Asp	Thr	Pro	Trp
			580					585					590		
lle	Ala	Arg	Thr	Leu	Asp	Asn	Leu	Ala	Asp	Glu	Leu	Thr	Ala	He	Leu
		595					600					605			
Ser	Ala	Pro	Ala	Lys	Leu	lle	Gly	His	Gly	Val	Lys	Gly	Val	Ser	Ser
	610					615					620				
Leu	Phe	Lys	Lys	llis	Lys	Leu	Pro	Phe							
625					630										

```
<211> 779
<212> PRT
<213> Homo sapiens
<400> 586
Met Leu Pro Cys His Ser Pro Ser Arg Ser Asp Gln Val Asn Leu Gly
 1
                  5
                                                          15
Pro Ser Ile Asn Ser Lys Leu Leu Gly Met Ser Thr Gln Asn Tyr Ala
             20
                                  25
Leu Met Gln Val Ala Gly Gln Glu Gly Thr Phe Ser Leu Val Ala Leu
                             40
Pro His Val Ala Ser Ala Gln Pro 11e Gln Lys Pro Arg Met Ser Leu
     50
                         55
                                              60
Pro Glu Asn Leu Lys Leu Pro 11e Pro Arg Tyr Gln Pro Pro Arg Asn
                     70
                                          75
Ser Lys Ala Ser Arg Lys Lys Pro Ile Leu Ile Phe Pro Lys Ser Gly
                 85
                                     90
Cys Ser Lys Ala Pro Ala Gln Thr Gln Met Cys Pro Gln Met Ser Pro
                                 105
                                                     110
Ser Pro Pro His His Pro Glu Leu Leu Tyr Lys Pro Ser Pro Phe Glu
                            120
Glu Val Pro Ser Leu Glu Gln Ala Pro Ala Ser Ile Ser Thr Ala Ala
    130
                        135
                                             140
Leu Thr Asn Gly Ser Asp His Gly Asp Leu Arg Pro Pro Val Thr Asn
145
                    150
                                         155
                                                             160
Thr His Gly Ser Leu Asn Pro Pro Ala Thr Pro Ala Ser Ser Thr Pro
                165
                                     170
Glu Glu Pro Ala Lys Gln Asp Leu Thr Ala Leu Ser Gly Lys Ala His
            180
                                 185
                                                     190
Phe Val Ser Lys Ile Thr Ser Ser Lys Pro Ser Ala Val Ala Ser Glu
                            200
                                                 205
Lys Phe Lys Glu Gln Val Asp Leu Ala Lys Thr Met Thr Asn Leu Ser
    210
                        215
                                             220
Pro Thr lle Leu Gly Asn Ala Val Gln Leu Ile Ser Ser Val Pro Lys
225
                    230
                                         235
                                                             240
```

Gly	Lys	Leu	Pro	lle	Pro	Pro	Tyr	Ser	Arg	Met	Lys	Thr	Met	Glu	Val
				245					250					255	
Tyr	Lys	He	Lys	Ser	Asp	Ala	Asn	lle	Ala	Gly	Phe	Ser	Leu	Pro	Gly
			260					265					270		
Pro	Lys	Ala	Asp	Cys	Asp	Lys	lle	Pro	Ser	Thr	Thr	Glu	Gly	Phe	Asn
		275					280					285			
Ala	Ala	Thr	Lys	Val	Ala	Ser	Arg	Leu	Pro	Val	Pro	Gln	Va]	Ser	Gln
	290					295					300				
Gln	Ser	Ala	Cys	Glu	Ser	Ala	Phe	Cys	Pro	Pro	Thr	Lys	Leu	Asp	Leu
305					310					315					320
Asn	His	Lys	Thr	Lys	Leu	Asn	Ser	Gly	Ala	Ala	Lys	Arg	Lys	G1 y	Arg
				325					330					335	
Lys	Arg	Lys	Val	Pro	Asp	Glu	11e	Leu	Ala	Phe	Gln	Gly	Lys	Arg	Arg
			340					345					350		
Lys	Tyr	He	lle	Asn	Lys	Cys	Arg	Asp	Gly	Lys	Glu	Arg	Val	Lys	Ásn
		355					360					365			
Asp	Pro	Gln	Glu	Phe	Arg	Asp	Gln	Lys	Leu	Gly	Thr	Leu	Lys	Lys	Tyr
	370					375					380				
Arg	Ser	lle	Met	Pro	Lys	Pro	He	Met	Val	He	Pro	Thr	Leu	Ala	Ser
385					390					395					400
Leu	Ala	Ser	Pro	Thr	Thr	Leu	Gln	Ser	Gln	Met	Leu	Gly	Gly	Leu	Gly
				405					410					415	
Gln	Asp	Val	Leu	Leu	Asn	Asn	Ser	Leu	Thr	Pro	Lys	Tyr	Leu	Gly	Cys
			420					425					430		
Lys	Gln	Asp	Asn	Ser	Ser	Ser	$P_{\mathbf{ro}}$	Lys	Pro	Ser	Ser	Val	Phe	Arg	Asn
		435					440					$44\widehat{5}$			
G1 y	Phe	Ser	Gly	Пе	Lys	Lys	Pro	Trp	His	Arg	Cys	His	Val	Cys	Asn
	450					455					460				
His	His	Phe	Gln	Phe	Lys	Gln	His	Leu	Arg	Asp	His	Met	Asn	Thr	His
465					470					475	•				480
Thr	Asn	Arg	Arg	Pro	Tyr	Ser	Cys	Arg	He	Cys	Arg	Lys	Ser	Tyr	Val
				485					490					495	
Arg	Pro	Gly	Ser	Leu	Ser	Thr	His	Met	Lys	Leu	His	His	Ġly	Glu	Asn
			500					505					510		
Arg	Leu	Lys	Lys	Leu	Met	Cys	Cys	Glu	Phe	Cys	Ala	Lys	Val	Phe	G1 y
		515					520					525			

His	lle	Arg	Val	Tyr	Phe	Gly	llis	Leu	Lys	Glu	Val	llis	Arg	Val	Val
	530					535					540				,
He	Ser	Thr	Glu	Pro	Ala	Pro	Ser	Glu	Leu	Gln	Pro	Gly	Λsp	He	Pro
545					550					555					560
Lys	Asn	Arg	Asp	Met	Ser	Val	Arg	Gly	Met	Glu	Gly	Ser	Leu	Glu	Arg
				565					570					575	
Glu	Asn	Lys	Ser	Asn	Leu	Glu	Glu	Asp	Phe	Leu	Leu	Asn	Gln	Ala	Asp
			580					585					590		
Glu	Val	Lys	Leu	Gln	lle	Lys	Cys	G1 y	Arg	Cys	Gln	Ile	Thr	Ala	Gln
		595					600					605			
Ser	Phe	Ala	Glu	Île	Lys	Phe	His	Leu	Leu	Gly	Val	His	Gly	Glu	Glu
	610					615					620				
He	Glu	Gly	Arg	Leu	Gln	Glu	Gly	Thr	Phe	Pro	Gly	Ser	Lys	Gly	Thr
625					630					635				٠	640
Gln	Glu	Glu	Leu	Va]	Gln	His	Ala	Ser	Pro	Asp	Trp	Lys	Arg	His	Pro
				645					650					655	
Glu	Arg	Gly	Lys	Pro	Glu	Lys	Val	His	Ser	Ser	Ser	Glu	Glu	Ser	His
			660					665					670		
Ala	Cys	Pro	Arg	Leu	Lys	Arg	Gln	Leu	His	Leu	llis	Gln	Asn	Gly	Val
		675					680					685			
Glu	Met	Leu	Met	Glu	Asn	Glu	Gly	Pro	Gln	Ser	Gly	Thr	Asn	Lys	Pro
	690					695					700				
Arg	Glu	Thr	Cys	Gln	Gly	Pro	Glu	Cys	Pro	Gly	Leu	His	Thr	Phe	Leu
705					710					715					720
Leu	Trp	Ser	His	Ser	Gly	Phe	Asn	Cys	Leu	Leu	Cys	Ala	Glu	Met	Leu
				725					730					735	
Gly	Arg	Lys	Glu	Asp	Leu	Leu	His	His	Trp	Lys	His	G1n	His	Asn	Cys
			740					745					750		
Glu	Asp	Pro	Ser	Lys	Leu	Trp	Ala	Пе	Leu	Asn	Thr	Val	Ser	Asn	Gln
		755					760					765			
Gly	Val	He	Glu	Leu	Ser	Ser	Glu	Ala	Glu	Lys					
	770					775									

<210> 4964

<211> 353

<212>	PRT		
<213>	Homo	sapiens	
<400>	587		

<400)> 58	37													
Met	Glu	Phe	Arg	Lys	Thr	Met	Asp	He	Asp	His	Thr	Leu	Asp	Trp	Gln
l				5					10					15	
Pro	Pro	Glu	Val	Ile	Gln	Lys	Tyr	Met	Pro	Gly	Gly	Leu	Cys	Gly	Tyr
			20					25					30		
Asp	Arg	Asp	Gly	Cys	Pro	Val	Trp	Tyr	Asp	Ile	Thr	Gly	Pro	Leu	Asp
		35					40					45			
Pro	Lys	Gly	Leu	Leu	Phe	Ser	Val	Thr	Lys	Gln	Asp	Leu	Leu	Lys	Thr
	50					55					60				
Lys	Met	Arg	Asp	Cys	Glu	Arg	He	Leu	His	Glu	Cys	Asp	Leu	Gln	Thr
65					70					75					80
Glu	Arg	Leu	Gly	Lys	Lys	He	Glu	Thr	He	Val	Met	He	Phe	Asp	Cys
				85					90					95	
Glu	Gly	Leu	Gly	Leu	Lys	His	Phe	Trp	Lys	Pro	Leu	Val	Glu	Val	Tyr
			100					105					110		
G1n	Glu	Phe	Phe	Gly	Leu	Leu	Glu	Glu	Asn	Tyr	Pro	Glu	Thr	Leu	Lys
		115					120					125			
Phe		Leu	lle	Val	Lys		Thr	Lys	Leu	Phe	Pro	Val	Gly	Tyr	Asn
	130					135					140				
	Met	Lys	Pro	Phe		Ser	Glu	Asp	Thr		Arg	Lys	He	He	
145				_	150					155			_		160
Leu	Gly	Asn	Asn		Lys	Glu	Gly	Leu		Lys	Leu	He	Ser		G] u
0.1		В		165	123	0.1	0.1	 .	170			• >		175	
Glu	Leu	Pro		GIn	Phe	Gly	Gly		Leu	Thir	Asp	Pro		Gly	Asn
		0	180	T)				185	0.1	0.1			190		
Pro	Lys		Leu	lhr	Lys	He	Asn	Tyr	GLy	G1 y	Glu		Pro	Lys	Ser
	т	195			C1		200	T)	<i>C</i> 1	ar.	<i>a</i> 1	205	C	,, ,	6.1
Met		val	Arg	Asp	61n		Lys	Ihr	GIn	lyr		liis	Ser	Val	GIn
	210		C.I.	C	C	215	C.1	12 1	6.1	ar.	220			121	Б
	Asn	Arg	61y	Ser		HIS	Gln	Val	Glu		Glu	He	Leu	Phe	
225	C	V = 1	1	Λ	230	Cl	DI	C .	C .	235	C1	A 1	۸.	1.3	240
01 A	cys	val	Leu		1.rp	GIN	Phe	ser		Asp	Gly	Ala	Asp		Gly
				245					250					255	

Phe Gly Val Phe Leu Lys Thr Lys Met Gly Glu Arg Gln Arg Ala Gly Glu Met Thr Glu Val Leu Pro Ser Gln Arg Tyr Asn Ala His Met Val Pro Glu Asp Gly Asn Leu Thr Cys Ser Glu Ala Gly Val Tyr Val Glu Ser Glu Ser Gly Lys Ser Cys Cys His Leu Pro Val Ile Ile Cys Ser His Glu Leu Gln Asn Ser His Ser Asn Ser Gln Val Met Ala Tyr Gln Met Val Arg Lys Cys Lys Leu Ser Arg Pro Leu Pro Leu Pro Ala Ser Asn

<210> 4965

<211> 606

<212> PRT

<213> Homo sapiens

<400> 588

Met Cys Leu Glu Lys Arg Tyr Leu Asp Val Leu Ser Asp Val Thr Gly Pro Gln Val Ser Cys Tyr Ile Thr Ala Pro Ser Tyr Val Leu Gln Gln Leu Glu Cys Arg Ile Ile Asn His Met Ser Ser Leu lle Val Gly Asp Asn Glu Glu Leu Val Ser Asn Val Ile Thr Ile Glu Cys Ser Asp Lys Glu Lys Arg Val Pro Phe Pro Ile Gly Ile Ala Ile Pro Phe Thr Ala Arg Tyr Arg Gly Asn Tyr Arg Asp lle Met Val Lys Val Cys Asp lle Asn Leu Gln Ser Ser Tyr Leu Asn Pro Asn Ser Leu Glu Gly Met Lys

Gly	Gly	Tyr 115	Lys	Gly	Thr	Cys	Ala 120	Ser	Val	Lys	Val	Tyr 125	Lys	Leu	Gly
lle		Ser	Val	Val	Ser		Leu	Lys	Lys	G1 u			Thr	Val	Thr
Lys	130 Lys	Gly	Leu	Ala	Leu	135 Lys	Ser	Ser	Met	Asp	140 Ser	Arg	lle	Ser	Leu
145					150					155					160
Asn	Tyr	Pro	Pro	Gly 165	Val	Phe	Thr	Ser	Pro 170	Val	Leu	Val	Gln	Leu 175	Lys
lle	Gln	Pro	Val 180	Asp	Pro	Ala	Leu	Val 185	Ala	His	Leu	Lys	Ala 190	Gln	Glm
Asp	Thr	Phe 195	Tyr	Ser	Val	Gln	Ser 200	Thr	Ser	Pro	Leu	Ile 205	His	Ile	Gln
His	Pro 210	Ser	Thr	Tyr	Pro	Phe 215	Gln	Lys	Pro	Val	Thr 220	Leu	Phe	Leu	Pro
Cys 225	Ser	Pro	Tyr	Leu	Asp 230	Lys	Asn	Asn	Leu	Gly 235	Ser	Glu	lle	Asp	His
Lys	Arg	Arg	Ala	Ser 245	Ala	Thr	lle	Asn	Arg 250	Ile	Thr	Pro	Ser	Tyr 255	Phe
Asn	Arg	Thr	Lys 260	Ile	Ala	Ser	Ile	Arg 265	Lys	Pro	Arg	Lys	Asn 270	Ala	Ser
Glu	Cys	Leu 275	Lys	Leu	Leu	Gly	Phe 280	Arg	Ser	Gln	Asp	Ser 285	Gly	Trp	Cys
Gly	Leu 290	Asp	Asp	Val	Val	Lys 295	Thr	He	Gln	Ser	Gly 300	Leu	Val	Ser	Val
G1u 305	Leu	Tyr	Glu	His	Leu 310	Glu	Arg	Phe	lle	Val 315	Leu	His	Leu	Ser	Ser 320
Thr	Met	Asp	Asn	Ser 325	His	Leu	Val	Thr	Phe 330	Val	Lys	Ser	Leu	Glu 335	Glu
Ala	Met	Leu	Ser 340	Thr	Thr	Ala	Cys	lle 345	Val	Leu	Ser	His	G1n 350	Lys	Asp
Asn	Pro	His 355	Arg	Ile	Ala	Val	Leu 360	Val	Val	Pro	Ser	Lys 365	Asp	Leu	Ser
Gln	Val 370	Leu	Lys	Asp	Leu	His 375	Leu	Glu	Gly	Phe	Gly 380	Gly	Pro	Pro	Glu
Pro	Ser	Arg	His	Phe	G1n	Val	Arg	G] u	Gly	Glu 205	Gln	Leu	Leu	Leu	Arg

Phe Thr Gly Asn Ile Phe Ala Ser Ser Asn Gly Lys Asp Tyr Gly Lys Asp Tyr Thr Leu Ile Phe His Leu Gln Arg Lys Pro Arg Leu Glu Leu Gln Ile Lys Glu Val Asp Glu Phe Gly Asn Tyr Ser Cys Pro His Tyr Lys Gly Thr Ile Val Val Tyr Lys Val Pro Lys Gly Lys Ile Val Pro Asn Leu Asn Gln Ser Leu Val Ile Asn Glu Asn His Ser Gln Leu Pro Ile Cys Lys Leu Pro Leu Lys Leu Pro Lys His Lys Lys Leu Ile Asn His Pro Gln Ser Thr Lys Arg Val Ser Lys Asp Pro Val Glu Ala Leu Trp Asp Asn Leu Leu His Trp Leu Ala Glu Glu Leu Ser Glu Glu Asn Ala Glu Ser Leu Ser Ser Thr Leu Pro Leu Arg Arg Ser Thr Ile Gln Leu Ile Lys Leu Lys Asn Pro Asp Asp Leu Thr Glu Gln Ile His Glu Phe Leu Cys Phe Trp Lys Lys Ser Leu Pro Thr Phe Thr Asp Lys Leu Arg Leu Leu Ala Arg His Leu Arg Lys lle Gly Arg Ser Asp Leu Ala Glu Glu Leu Lys Phe Lys Trp Glu Asn Lys Val Phe Thr Glu

<210> 4966

<211> 151

<212> PRT

<213> Homo sapiens

<400> 589

Met Ala Thr Ile Thr Met Gln Ala Tyr Ser Arg Gly Phe Leu Ala Arg

1 5 10 15

Arg Arg Tyr Arg Lys Met Leu Glu Glu His Lys Ala Val Ile Leu Gln 20 25 Lys Tyr Ala Arg Ala Trp Leu Ala Arg Arg Arg Phe Gln Ser 11e Arg 35 40 45 Arg Phe Val Leu Asn Ile Gln Leu Thr Tyr Arg Val Gln Arg Leu Gln 55 Lys Lys Leu Glu Asp Gln Asn Lys Glu Asn His Gly Leu Val Glu Lys 65 70 75 80 Leu Thr Ser Leu Ala Ala Leu Arg Ala Gly Asp Val Glu Lys Ile Gln 85 90 Lys Leu Glu Ala Glu Leu Glu Lys Ala Ala Thr His Arg Arg Asn Tyr 100 105 Glu Glu Lys Gly Lys Arg Tyr Arg Asp Ala Val Glu Glu Val Ser Cys 115 120 125 Gly Arg Val Arg Arg Asp Gly Asn Met Arg Thr Leu Cys Glu Lys Asp 130 135 140 Glu Arg Cys Gln Leu Phe Leu 145 150

<210> 4967

<211> 731

<212> PRT

<213> Homo sapiens

<400> 590

Met Met Gln Ala Gln Glu Ser Leu Thr Leu Glu Asp Val Ala Val Asp 1 5 10 15

Phe Thr Trp Glu Glu Trp Gln Phe Leu Ser Pro Ala Gln Lys Asp Leu
20 25 30

Tyr Arg Asp Val Met Leu Glu Asn Tyr Ser Asn Leu Val Ala Val Gly
35 40 45

Tyr Gln Ala Ser Lys Pro Asp Ala Leu Ser Lys Leu Glu Arg Gly Glu 50 55 60

Glu Thr Cys Thr Thr Glu Asp Glu 11e Tyr Ser Arg 11e Cys Ser Glu 65 70 75 80

lle	Arg	Lys	He	Asp	Asp	Pro	Leu	Gln	His	His	Leu	Gln	Asn	G1n	Ser
				85					90					95	
11e	Gln	Lys	Ser	Val	Lys	Gln	Cys	His	Glu	Gln	Asn	Met	Phe	Gly	Asn
			100					105					110		
lle	Val	Asn	Gln	Asn	Lys	Gly	His	Phe	Leu	Leu	Lys	Gln	Asp	Cys	Asp
		115					120					125			
Thr	Phe	Asp	Leu	His	Glu	Lys	Pro	Leu	Lys	Ser	Asn	Leu	Ser	Phe	Glu
	130					135					140				
Asn	Gln	Lys	Arg	Ser	Ser	Gly	Leu	Lys	Asn	Ser	Ala	Glu	Phe	Asn	Arg
145					150					155					160
Asp	Gly	Lys	Ser	Leu	Phe	His	Ala	Asn	His	Lys	Gln	Phe	Tyr	Thr	Glu
				165					170					175	
Met	Lys	Phe	Pro	Ala	He	Ala	Lys	Pro	lle	Asn	Lys	Ser	Gln	Phe	lle
			180					185					190		
Lys	Gln	Gln	Arg	Thr	His	Asn	He	Glu	Asn	Ala	His	Val	Cys	Ser	Glu
		195					200					205			
Cys	Gly	Lys	Ala	Phe	Leu	Lys	Leu	Ser	Gln	Phe	Ile	Asp	His	Gln	Arg
	210					215					220				
Val	His	Thr	Gly	Glu	Lys	Pro	His	Val	Cys	Ser	Met	Cys	Gly	Lys	Ala
225					230					235					240
Phe	Ser	Arg	Lys	Ser	Arg	Leu	Met	Asp	His	Gln	Arg	Thr	His	Thr	Glu
				245					250					255	
Leu	Lys	His	Tyr	Glu	Cys	Thr	Glu	Cys	Asp	Lys	Thr	Phe	Leu	Lys	Lys
			260					265					270		
Ser	Gln	Leu	Asn	He	His	Gln	Lys	Thr	His	Met	Gly	Gly	Lys	Pro	Tyr
		275					280					285			
Thr		Ser	Gln	Cys	Gly		Ala	Phe	lle	Lys	Lys	Cys	Arg	Leu	He
	290					295					300				
Tyr	His	Gln	Arg	Thr		Thr	Gly	Glu	Lys		His	Gly	Cys	Ser	
305					310					315					320
Cys	Gly	Lys	Ala		Ser	Thr	Lys	Phe		Leu	Thr	Thr	His	Gln	Lys
				325				,	330					335	
Thr	His	Thr		Glu	Lys	Pro	Tyr		Cys	Ser	Glu	Cys		Lys	Gly
			340			_		345					350	m.	
Phe	He		Lys	Arg	Arg	Leu		Ala	His	His	Arg		His	Thr	Gly
		355					360					365			

Glu		Pro	Phe	He	Cys		Lys	Cys	Gly	Lys		Phe	Thr	Leu	Lys
	370					375					380				
Asn 385	Ser	Leu	lle	Thr	His 390	Gln	Gln	Thr	His		Gly	Glu	Lys	Leu	
		C	C1				C1	131	0	395			0		400
ınr	Cys	Ser	61 u		Gly	Lys	GIy	Phe		Met	Lys	HIS	Cys		Met
				405					410					415	
Val	His	GIn		Thr	His	Thr	Gl y	G] u	Lys	Pro	Tyr	Lys	Cys	Asn	Glu
			420					425					430		
Cys	Gly	Lys	Gly	Phe	Ala	Leu		Ser	Pro	Leu	Ile	Arg	His	Gln	Arg
		435					440					445			
Thr	His	Thr	Gly	Glu	Lys	Pro	Tyr	Val	Cys	Thr	Glu	Cys	Arg	Lys	Gly
	450					455					460				
Phe	Thr	Met	Lys	Ser	Asp	Leu	lle	Val	His	Gln	Arg	Thr	His	Thr	Ala
465					470					475					480
Glu	Lys	Pro	Tyr	lle	Cys	Asn	Asp	Cys	Gly	Lys	Gly	Phe	Thr	Val	Lys
				485					490					495	
Ser	Arg	Leu	lle	Val	His	Gln	Arg	Thr	His	Thr	Gly	Glu	Lys	Pro	Tyr
			500					505					510		
Val	Cys	Gly	Glu	Cys	Gly	Lys	Gly	Phe	Pro	Ala	Lys	Πle	Arg	Leu	Met
		515					520					525			
Gly	His	Gln	Arg	Thr	His	Thr	Gly	Glu	Lys	Pro	Tyr	Ile	Cys	Asn	Glu
	530					535					540				
Cys	Gly	Lys	Gly	Phe	Thr	Glu	Lys	Ser	His	Leu	Asn	Val	His	Arg	Arg
545					550					555					560
Thr	His	Thr	Gly	Glu	Lys	Pro	Tyr	Val	Cys	Ser	Glu	Cys	Gly	Lys	Asp
				565					570					575	
Leu	Thr	Gly	Lys	Ser	Met	Leu	He	Ala	His	Gln	Arg	Thr	His	Thr	Gly
			580					585					590		
Glu	Lys	Pro	Tyr	He	Cys	Asn	Glu	Cys	Gly	Lys	Gly	Phe	Thr	Met	Lys
		595					600					605			
Ser	Thr	Leu	Ser	lle	His	Gln	Gln	Thr	His	Thr	Gly	Glu	Lys	Pro	Tyr
	610					615					620				
Lys	Cys	Asn	Glu	Cys	Asp	Lys	Thr	Phe	Arg	Lys	Lys	Thr	Cys	Leu	lle
625					630					635					640
Gle	Hic	Gla	Ara	Pho	Hic	Thr	C1v	Lve	The	Sar	Dho	A1.c	Cvc	The	C1

Cys Gly Lys Phe Ser Leu Arg Lys Asn Asp Leu Ile Thr His Gln Arg Ile His Thr Gly Glu Lys Pro Tyr Lys Cys Ser Asp Cys Gly Lys Ala Phe Thr Thr Lys Ser Gly Leu Asn Val His Gln Arg Lys His Thr Gly Glu Arg Pro Tyr Gly Cys Ser Asp Cys Gly Lys Ala Phe Ala His Leu Ser Ile Leu Val Lys His Arg Arg Ile His Arg

<210> 4968

<211> 1186

<212> PRT

<213> Homo sapiens

<400> 591

Met Tyr Gln Ala Leu Glu Gln Gly Gly Met Thr Phe Gly Trp Val Cys Trp Met Ile Leu Phe Asp Ser Ser Leu Tyr Phe Leu Cys Gly Trp Tyr Leu Ser Asn Leu lle Pro Gly Thr Phe Gly Leu Arg Lys Pro Trp Tyr Phe Pro Phe Thr Ala Ser Tyr Trp Lys Ser Val Gly Phe Leu Val Glu Lys Arg Gln Tyr Phe Leu Ser Ser Ser Leu Phe Phe Phe Asn Glu Asn Phe Asp Asn Lys Gly Ser Ser Leu Gln Asn Arg Glu Gly Glu Leu Glu Gly Ser Ala Pro Gly Val Thr Leu Val Ser Val Thr Lys Glu Tyr Glu Gly His Lys Ala Val Val Gln Asp Leu Ser Leu Thr Phe Tyr Arg Asp

Gln Ile Thr Ala Leu Leu Gly Thr Asn Gly Ala Gly Lys Thr Thr 11e

	130	1				135					140				
Īle	Ser	Met	Leu	Thr	Gly	Leu	His	Pro	Pro	Thr	Ser	Gly	Thr	He	He
145					150					155	!				160
lle	Asn	Gly	Lys	Asn	Leu	Gln	Thr	Asp	Leu	Ser	Arg	Val	Arg	Met	Glu
				165					170					175	
Leu	Gly	Val	Cys	Pro	Gln	Gln	Asp	lle	Leu	Leu	Asp	Asn	Leu	Thr	Val
			180					185					190		
Arg	Glu	His	Leu	Leu	Leu	Phe	Ala	Ser	lle	Lys	Ala	Pro	Gln	Trp	Thr
		195					200					205			
Lys	Lys	Glu	Leu	His	Gln	Gln	Val	Asn	Gln	Thr	Leu	Gln	Asp	Val	Asp
	210					215					220				
Leu	Thr	Gln	His	Gln	His	Lys	Gln	Thr	Arg	Ala	Leu	Ser	Gly	G1 y	Leu
225					230					235					240
Lys	Arg	Lys	Leu	Ser	Leu	Gly	He	Ala	Phe	Met	Gly	Met	Ser	Arg	Thr
				245					250					255	
Val	Val	Leu	Asp	Glu	Pro	·Thr	Ser	G1 y	Val	Asp	Pro	Cys	Ser	Arg	His
			260					265					270		
Ser	Leu	Trp	Asp	He	Leu	Leu	Lys	Tyr	Arg	Glu	Gly	Arg	Thr	He	lle
		275					280					285			
Phe		Thr	His	His	Leu	Asp	Glu	Ala	Glu	Ala	Leu	Ser	Asp	Arg	Val
	290					295					300				
	Val	Leu	GIn	His		Arg	Leu	Arg	Cys	Cys	G1 y	Pro	Pro	Phe	Cys
305		0.7			310					315					320
Leu	Lys	Glu	Ala	Tyr	Gly	Gln	G1 y	Leu		Leu	Thr	Leu	Thr		G1n
D	0	v 3	•	325					330					335	
Pro	Ser	Val		Glu	Ala	His	Asp		Lys	Asp	Met	Ala		Val	Thr
C		7.1	340	7.1	T		Б	345	. 1	D)		_	350		
Ser	Leu		Lys	He	lyr	He		GIn	Ala	Phe	Leu		Asp	Ser	Ser
C1	C	355	1	ть	т	TI	360	D	,		T)	365			
GIŸ		GIU	Leu	Thr	lyr		11e	Pro	Lys	Asp		Asp	Lys	Ala	Cys
Lou	370	Clv	Lou	Dho	C15	375	1	Λ	C1	Λ	380		C1		
385	LyS	оту	Leu	Phe	390	АТА	reu	ASP	GIY		Leu	HIS	Gin	Leu	
	Thr	G1v	Tvr	Gly		Sor	Acn	Thr	Thr	395	C1	Class	V _C 1	Dha	400
Jeu	1113	01 y	1 9 1	405	116	261	ush	1111	410	rea	oru	oru	val	415	Leu
Met	Leu	Leu	Gln	Asp	Ser	Asn	Lve	lve		Hie	116	Ala	ا ما		Thr
					~ ~ .	1 1			-		3 a C	4 1 1 CI	1. U U	O 1 Y	1 1 1 1

			420					425					430		
Glu	Ser	Glu	Leu	Gln	Asn	His	Arg	Pro	Thr	Gly	His	Leu	Ser	Gly	Tyr
		435					440					445			
Cys	Gly	Ser	Leu	Ala	Arg	Pro	Ala	Thr	Val	Gln	Gly	Val	Gln	Leu	Leu
	450					455					460				
Arg	Ala	Gln	Val	Ala	Ala	He	Leu	Ala	Arg	Arg	Leu	Arg	Arg	Thr	Leu
465					470					475					480
Arg	Ala	Gly	Lys	Ser	Thr	Leu	Ala	Asp	Leu	Leu	Leu	Pro	Val	Leu	Phe
				485					490					495	
Val	Ala	Leu	Ala	Met	Gly	Leu	Phe	Met	Val	Arg	Pro	Leu	Ala	Thr	Glu
			500					505					510		
Tyr	Pro	Pro	Leu	Arg	Leu	Thr	Pro	Gly	His	Tyr	Gln	Arg	Ala	Glu	Thr
		515					520					525			
Tyr	Phe	Phe	Ser	Ser	Gly	Gly	Asp	Asn	Leu	Asp	Leu	Thr	Arg	Val	Leu
	530					535					540				
Leu	Arg	Lys	Phe	Arg	Asp	Gln	Asp	Leu	Pro	Cys	Ala	Asp	Leu	Asn	Pro
545					550					555					560
Arg	Gln	Lys	Asn	Ser	Ser	Cys	Trp	Arg	Thr	Asp	Pro	Phe	Ser	His	Pro
				565					570					575	
Glu	Phe	Gln	Asp	Ser	Cys	Gly	Cys	Leu	Lys	Arg	Pro	Asn	Arg	Ser	Ala
			580					585					590		
Ser	Ala	Pro	Tyr	Leu	Thr	Asn	His	Leu	Gly	His	Thr	Leu	Leu	Asn	Leu
		595					600					605			
Ser	G1 y	Phe	Asn	Met	Glu	Glu	Tyr	Leu	Leu	Ala	Pro	Ser	Glu	Lys	Pro
	610					615					620				
Arg	Leu	Gly	Gly	Trp	Ser	Phe	Gly	Leu	Lys	Ile	Pro	Ser	Glu	Ala	Gly
625					630					635					640
Gly	Ala	Asn	Gly	Asn	He	Ser	Lys	Pro	Pro	Thr	Leu	Ala	Lys	Val	Trp
				645					650					655	
Tyr	Asn	Gln	Lys	Gly	Phe	His	Ser	Leu	Pro	Ser	Tyr	Leu	Asn	His	Leu
			660					665					670		
Asn	Asn	Leu	lle	Leu	Trp	G]n	His	Leu	Pro	Pro	Thr	Val	Asp	Trp	Arg
		675					680					685			
Gln		Gly	He	Thr	Leu		Ser	His	Pro	Tyr	Gly	Gly	Ala	Leu	Leu
	690					695					700				
Asn	Glu	Asp	Lys	He	Leu	Glu	Ser	He	Arg	Gln	Cvs	G1 v	Val	Ala	Leu

705					710					715					720
Cys	lle	Val	Leu	Gly	Phe	Ser	Ile	Leu	Ser	Ala	Ser	He	Gly	Ser	Ser
	,			725					730					735	
Val	Val	Arg	Asp	Arg	Val	Ile	Gly	Ala	Lys	Arg	Leu	Gln	His	lle	Ser
			740					745					750		
Gly	Leu	Gly	Tyr	Arg	Met	Tyr	Trp	Phe	Thr	Asn	Phe	Leu	Tyr	Asp	Met
		755					760					765			
Leu	Phe	Tyr	Leu	Val	Ser	Val	Cys	Leu	Cys	Val	Ala	Val	lle	Val	Ala
	770					775					780				
Phe	Gln	Leu	Thr	Ala	Phe	Thr	Phe	Arg	Lys	Asn	Leu	Ala	Ala	Thr	Ala
785					790					795					800
Leu	Leu	Leu	Ser	Leu	Phe	Gly	Tyr	Ala	Thr	Leu	Pro	Trp	Met	Tyr	Leu
				805					810					815	
Met	Ser	Arg	He	Phe	Ser	Ser	Ser	Asp	Val	Ala	Phe	He	Ser	Tyr	Val
			820					825					830		
Ser	Leu	Asn	Phe	He	Phe	Gly	Leu	Cys	Thr	Met	Leu	lle	Thr	He	Met
		835					840					845			
Pro	Arg	Leu	Leu	Ala	Пе	lle	Ser	Lys	Ala	Lys	Asn	Leu	Gln	Asn	Ile
	850					855					860				
Tyr	Asp	Val	Leu	Lys	Trp	Val	Phe	Thr	Ile	Phe	Pro	Gln	Phe	Cys	Leu
865					870					875					880
Gly	Gln	Gly	Leu	Val	Glu	Leu	Cys	Tyr	Asn	Gln	Thr	Lys	Tyr	Asp	Leu
				885					890					895	
Thr	His	Asn	Phe	Gly	He	Asp	Ser	Tyr	Val	Ser	Pro	Phe	Glu	Met	Asn
			900					905					910		
Phe	Leu	Gly	Trp	He	Phe	Val	Gln	Leu	Ala	Ser	Gln		Thr	Val	Leu
		915					920					925			
Leu		Leu	Arg	Val	Leu	Leu	His	Trp	Asp	Leu		Arg	Trp	Pro	Arg
	930					935					940				
	His	Ser	Thr	Leu		Gly	Thr	Val	Lys		Ser	Lys	Asp	Thr	
945					950					955					960
Val	Glu	Lys	GJu		Lys	Arg	Val	Phe		Gly	Arg	Thr	Asn		Asp
				965			_		970	-				975	
ile	Leu	Val		fyr	Asn	Leu	Ser		His	Tyr	Arg	Arg		Phe	Gln
	.,	7.7	980		0.1		T 1	985		0.1		n	990	0.1	
Acn	110		Ala	V a l	Gln	Acn	IΙο	Sor	Lou	CLv	110	Dro	1 40	$C \perp U$	$C I \cdots$

Cys Phe Gly Leu Leu Gly Val Asn Gly Ala Gly Lys Ser Thr Thr Phe Lys Met Leu Asn Gly Glu Val Ser Leu Thr Ser Gly His Ala Ile lle Arg Thr Pro Met Gly Asp Ala Val Asp Leu Ser Ser Ala Gly Thr Ala Gly Val Leu Ile Gly Tyr Cys Pro Gln Gln Asp Ala Leu Asp Glu Leu Leu Thr Gly Trp Glu His Leu Tyr Tyr Tyr Cys Ser Leu Arg Gly Ile Pro Arg Gln Cys lle Pro Glu Val Ala Gly Asp Leu Ile Arg Arg Leu His Leu Glu Ala His Ala Asp Lys Pro Val Ala Thr Tyr Ser Glv Glv Thr Lys Arg Lys Leu Ser Thr Ala Leu Ala Leu Val Gly Lys Pro Asp lle Leu Leu Leu Asp Glu Pro Ser Ser Gly Met Asp Pro Cys Ser Lys Arg Tyr Leu Trp Gln Thr lle Met Lys Glu Val Arg Glu Gly Cys Ala Ala Val Leu Thr Ser His Arg Phe Trp Tyr Gln Asp Asp Ala Gly Leu - 1175 He Lys <210> 4969 <211> 140 <212> PRT <213> Homo sapiens <400> 592 Met Tyr 11e Phe Phe Phe Phe Leu Lys Gln Met Ser Phe Asp Pro Asn l Leu Leu His Asn Asn Gly His Asn Gly Tyr Pro Asn Gly Thr Ser Ala

Ala Leu Arg Glu Thr Gly Val 11e Glu Lys Leu Leu Thr Ser Tyr Gly Phe lle Gln Cys Ser Glu Arg Gln Ala Arg Leu Phe Phe His Cys Ser Gln Tyr Asn Gly Asn Leu Gln Asp Leu Lys Val Gly Gly Asn Leu Ser Val Leu Leu Cys Lys Asn Val Ile Thr Lys Phe Val Leu His Ile Asn Leu Phe Met Ser Val Phe Gln Lys Val Leu Cys Lys Leu Lys His Phe Gly Leu Leu Val Val Leu Phe Ser Glu Glu Tyr Leu Tyr Lys Pro Ala Glu Ser Leu Lys Asp Tyr 11e Cys 11e Tyr Leu Gln

<210> 4970

<211> 177

<212> PRT

<213> Homo sapiens

<400> 593

Met Ser Ser Glu Thr Pro Thr Ser Arg Gln Leu Ser Glu Tyr Leu Lys His Ala Lys Gly Arg Thr Arg Thr Ala lle Arg Asn Gly Gln Val Trp Glu Glu Ser Leu Lys Arg Leu Arg Gln Lys Ala Ser Leu Thr Asn Val Thr Asp Pro Ser Leu Asp Leu Thr Ser Leu Ser Leu Glu Val Gly Cys Gly Ala Pro Ala Pro Val Val Arg Cys Asp Pro Cys Ser Pro Tyr Arg Thr lle Thr Gly Asp Cys Asn Asn Arg Trp Arg Gly Leu Gly Cys Gly

Gly Arg Pro Phe Gln Pro Leu Arg Pro Ala Leu Pro Arg Pro Leu Ser

<210> 4971

<211> 930

<212> PRT

<213> Homo sapiens

<400> 594

 Met
 Ala
 Gly
 Ala
 A

Asn Arg Phe Met 11e Glu Cys Asp 11e Cys Lys Asp Trp Phe His Gly 55 Ser Cys Val Gly Val Glu Glu His His Ala Val Asp Ile Asp Leu Tyr 70 65 75 80 His Cys Pro Asn Cys Ala Val Leu His Gly Ser Ser Leu Met Lys Lys 85 90 Arg Arg Asn Trp His Arg His Asp Tyr Thr Glu Ile Asp Asp Gly Ser 105 110 Lys Pro Val Gln Ala Gly Thr Arg Thr Phe 11e Lys Glu Leu Arg Ser 115 120 125

Arg Val Phe Pro Ser Ala Asp Glu lle lle lle Lys Met His Gly Ser

	130					135					140				
GIn	Leu	Thr	Gln	Arg	Tyr	Leu	Glu	Lys	His	Gly	Phe	Asp	Val	Pro	Πle
145					150					155					160
Met	Val	Pro	Lys	Leu	Asp	Asp	Leu	Gly	Leu	Arg	Leu	Pro	Ser	Pro	Thr
				165					170					175	
Phe	Ser	Val	Met	Asp	Val	Glu	Arg	Tyr	Val	Gly	Gly	Asp	Lys	Val	Ile
			180					185					190		
Asp	Val	He	Asp	Val	Ala	Arg	Gln	Ala	Asp	Ser	Lys	Met	Thr	Leu	His
		195					200					205			
Asn	Tyr	Val	Lys	Tyr	Phe	Met	Asn	Pro	Asn	Arg	Pro	Lys	Val	Leu	Asn
	210					215					220				
Val	Ile	Ser	Leu	Glu	Phe	Ser	Asp	Thr	Lys	Met	Ser	Glu	Leu	Val	Glu
225					230					235					240
Val	Pro	Asp	lle	Ala	Lys	Lys	Leu	Ser	Trp	Va]	Glu	Asn	Tyr	Trp	Pro
				245					250					255	
Asp	Asp	Ser	Val	Phe	Pro	Lys	Pro	Phe	Val	Gln	Lys	Tyr	Cys	Leu	Met
			260					265					270		
Gly	Val	Gln	Asp	Ser	Tyr	Thr	Asp	Phe	His	He	Asp	Phe	Gly	Gly	Thr
		275					280					285			
Ser	Val	Trp	Tyr	His	Val	Leu	Trp	Gly	Glu	Lys	He	Phe	Tyr	Leu	He
	290					295					300				
Lys	Pro	Thr	Asp	Glu	Asn	Leu	Ala	Arg	Tyr	Glu	Ser	Trp	Ser	Ser	
305					310					315					320
Val	Thr	Gln	Ser	Glu	Val	Phe	Phe	Gly	Asp	Lys	Val	Asp	Lys	Cys	Tyr
			•	325					330					335	
Lys	Cys	Val		Lys	Gln	Gly	His		Leu	Phe	Val	Pro		Gly	Trp
			340					345					350		
He	His		Val	Leu	Thr	Ser		Asp	Cys	Met	Ala		G] y	Gly	Asn
	_	355		_			360					365	_		
Phe		His	Asn	Leu	Asn	lle	Gly	Met	GIn	Leu		Cys	Tyr	Glu	Met
0.1	370		,	,	7 73	375		,	D.		380	ъ.	int	131	0.1
	Lys	Arg	Leu	Lys		Pro	Asp	Leu	Phe		Phe	Pro	Phe	Phe	
385	2.1	C	Tr.	121	390	. 1	,			395	<i>(</i>) 1	TI	,	,	400
Ala	116	Cys	rp		val	Ala	Lys	Asn		Leu	61u	Inr	Leu		ыu
1	Λ 33 -	C1	Λ ~	405	Dh -	Gln	D~ -	C1	410	Т	1	V ₀ 1	C1.	415	V e. 1
1 5 11	MY	11111	ΔSD	UIV	E LIE	(111)	1. (1)	(111)	1111	1 1/1	1.01	V CLI	OTH	GIV	1 71 1

				420					425					430		
	Lys	Ala	Leu	His	Thr	Ala	Leu	Lys	Leu	Trp	Met	Lys	Lys	Glu	Leu	Val
			435					440					445			
	Ser	Glu	His	Ala	Phe	Glu	lle	Pro	Лѕр	Asn	Val	Arg	Pro	Gly	His	Leu
		450					455					460				
•	He	Lys	Glu	Leu	Ser	Lys	Val	Пe	Arg	Ala	He	Glu	Glu	Glu	Asn	Gly
	465					470					475					480
	Lys	Pro	Val	Lys	Ser	Gln	Gly	11e	Pro	lle	Val	Cys	Pro	Val	Ser	Arg
					485					490					495	
	Ser	Ser	Asn	Glu	Ala	Thr	Ser	Pro	Tyr	His	Ser	Arg	Arg	Lys	Met	Arg
				500					505					510		
	Lys	Leu	Arg	Asp	His	Asn	Val	Arg	Thr	Pro	Ser	Asn	Leu	Asp	11e	Leu
			515					520					525			
	Glu	Leu	His	Thr	Arg	Glu	Val	Leu	Lys	Arg	Leu	Glu	Met	Cys	Pro	Trp
		530					535					540				
	Glu	Glu	Asp	He	Leu	Ser	Ser	Lys	Leu	Asn	Gly	Lys	Phe	Asn	Lys	His
	545					550					555					560
	Leu	Gln	Pro	Ser	Ser	Thr	Val	Pro	Glu	Trp	Arg	Ala	Lys	Asp	Asn	Asp
					565					570					575	
	Leu	Arg	Leu	Leu	Leu	Thr	Asn	Gly	Arg	He	He	Lys	Asp	Glu	Arg	Gln
				580					585					590		
	Pro	Phe	Ala	Asp	Gln	Ser	Leu	Tyr	Thr	Ala	Asp	Ser	Glu	Asn	Glu	Glu
			595					600					605			
	Asp		Arg	Arg	Thr	Lys	Lys	Ala	Lys	Met	Lys		Glu	Glu	Ser	Ser
		610					615					620				
		Val	Glu	Gly	Val		His	Glu	Glu	Ser		Lys	Pro	Leu	Asn	
	625					630					635					640
	Phe	Phe	Thr	Arg		Lys	Ser	Glu	Leu		Ser	Arg	Ser	Ser		Tyr
					645		0.1			650	D		_	4 2.1	655	,
	Ser	Asp	He		Glu	Ser	Glu	Asp		Gly	Pro	Glu	Cys		Ala	Leu
		C	1.1	660	TI	TI.	C1	C1	665	C1	C	C .	C I	670	C1	1
	Lys	Ser		Phe	Ihr	inr	Glu		Ser	Glu	Ser	Ser		Asp	GIU	Lys
	1.	C1 :	675	11.	TI	C.	Λ.	680	1.	C1	. С 1	C	685	V. 1	M. 4	Λ.
	Lys		G1U	116	ınr	ser	Asn	rne	Lys	GIU	61 u		ASN	val	мет	arg
	Acn	690	Lou	Cln	1	S	695	Lva	Dna	Sar	A 20.00	700 San	C1	11.	Dag	71,

705					710					715					720
Lys	Arg	Glu	Cys	Pro	Thr	Ser	Thr	Ser	Thr	Glu	Glu	Glu	Ala	He	Gln
				725					730					735	
Gly	Met	Leu	Ser	Met	Ala	Gly	Leu	His	Tyr	Ser	Thr	Cys	Leu	Gln	Arg
			740					745					750		
Gln	Ile	Gln	Ser	Thr	Asp	Cys	Ser	G1 y	Glu	Arg	Asn	Ser	Leu	Gln	Asp
		755					760					765			
Pro	Ser	Ser	Cys	His	Gly	Ser	Asn	His	Glu	Val	Λrg	Gln	Leu	Tyr	Arg
	770					775					780				
Tyr	Asp	Lys	Pro	Val	Glu	Cys	Gly	Tyr	His	Val	Lys	Thr	Glu	Asp	Pro
785					790					795					800
Asp	Leu	Arg	Thr	Ser	Ser	Trp	He	Lys	Gln	Phe	Asp	Thr	Ser	Arg	Phe
				805					810					815	
His	Pro	Gln	Asp	Leu	Ser	Arg	Ser	Gln	Lys	Cys	lle	Arg	Lys	Glu	Gly
			820					825					830		
Ser	Ser	Glu	lle	Ser	Gln	Arg	Val	Gln	Ser	Arg	Asn	Tyr	Val	Asp	Ser
		835					840					845			
Ser	Gly	Ser	Ser	Leu	Gln	Asn	Gly	Lys	Tyr	Met	Gln	Asn	Ser	Asn	Leu
	850					855					860				
Thr	Ser	Gly	Ala	Cys	Gln	Ile	Ser	Asn	Gly	Ser	Leu	Ser	Pro	Glu	Arg
865					870					875					880
Pro	Val	Gly	Glu	Thr	Ser	Phe	Ser	Val	Pro	Leu	His	Pro	Thr	Lys	Arg
				885					890					895	
Pro	Ala	Ser	Asn	Pro	His	Leu	Ser	Λla	Thr	Arg	Gln	G]n	Lys	Val	Asn
			900					905					910		
Val	Gln	Lys	Lys	Glu	Trp	Gln	Gln	Pro	Asn	Asn	Val	Leu	G1y	Arg	Ser
		915					920					925			
Leu															
	930														

<210> 4972

<211> 755

<212> PRT

<213> Homo sapiens

<400)> 59	95													
Met	Arg	Lys	Val	Lys	Lys	Leu	Arg	Leu	Asp	Lys	Glu	Asn	Thr	Gly	Ser
1				5					10					15	
Trp	Arg	Ser	Phe	Ser	Leu	Asn	Ser	Glu	Gly	Ala	Glu	Arg	Met	Ala	Thr
			20					25					30		
Ser	Tyr	Asp	Phe	His	Ser	Glu	Ser	Gly	Leu	Phe	Leu	Phe	Gln	Ala	Ser
		35					40					45			
Asn	Ser	Leu	Phe	His	Cys	Arg	Asp	Gly	Gly	Lys	Asn	Gly	Phe	Met	Val
	50					55					60				
Ser	Pro	Met	Lys	Pro	Leu	Glu	Ile	Lys	Thr	Gln	Cys	Ser	Gly	Pro	Arg
65					70					75					80
Met	Asp	Pro	Lys	Ile	Cys	Pro	Ala	Asp	Pro	Ala	Phe	Phe	Ser	Phe	He
				85					90					95	
Asn	Asn	Ser	Asp	Leu	Trp	Val	Ala	Asn	He	Glu	Thr	Gly	Glu	Glu	Arg
			100					105					110		
Arg	Leu	Thr	Phe	Cys	His	Gln	Gly	Leu	Ser	Asn	Val	Leu	Asp	Asp	Pro
		115					120					125			
Lys		Ala	Gly	Val	Ala		Phe	Val	Ile	Gln	Glu	Glu	Phe	Asp	Arg
	130					135					140				
Phe	Thr	Gly	Tyr	Trp	Trp	Cys	Pro	Thr	Ala	Ser	Trp	Glu	Gly	Ser	Glu
145					150					155					160
Gly	Leu	Lys	Thr		Arg	He	Leu	Tyr		Glu	Val	Asp	Glu		Glu
				165					170					175	
Val	Glu	Val		His	Val	Pro	Ser		Ala	Leu	Glu	Glu		Lys	Thr
	_		180	_				185	_				190		
Asp	Ser		Arg	Tyr	Pro	Arg		Gly	Ser	Lys	Asn		Lys	He	Ala
		195		6.1	Б.		200		_			205			
Leu		Leu	Ala	Glu	Phe		Thr	Asp	Ser	GIn	Gly	Lys	He	Val	Ser
<i>m</i> 1	210	0.1				215			-		220		153	Б	
	GIn	Glu	Lys	Glu		Val	GIn	Pro	Phe		Ser	Leu	Phe	Pro	
225	61	T	7.1	. 1	230	4.1	0.1	Tr.	T)	235		61	,	T	240
Val	Glu	lyr	He		Arg	Ala	Gly	Trp		Arg	Asp	GIy	Lys		Ala
æ	4 3		DI	245			ь	63	250	т		63		255	
lrp	Ala	Met		Leu	Asp	Arg	Pro		GIn	Trp	Leu	GIn		val	Leu
			260					265					270		

Leu	Pro	Pro	Ala	Leu	Phe	Ile	Pro	Ser	Thr	Glu	Asn	Glu	Glu	Gln	Arg
		275					280					285			
Leu	Λla	Ser	Ala	Arg	Ala	Val	Pro	Arg	Asn	Val	Gln	Pro	Tyr	Val	Val
	290					295					300				
Tyr	Glu	Glu	Val	Thr	Λsn	Val	Trp	He	Asn	Val	His	Asp	Ile	Phe	Tyr
305					310					315					320
Pro	Phe	Pro	Gln	Ser	Glu	Gly	Glu	Asp	Glu	Leu	Cys	Phe	Leu	Arg	Ala
				325					330					335	
Asn	Glu	Cys	Lys	Thr	Gly	Phe	Cys	His	Leu	Tyr	Lys	Val	Thr	Ala	Val
			340					345					350		
Leu	Lys	Ser	G1n	Gly	Tyr	Asp	Trp	Ser	Glu	Pro	Phe	Ser	Pro	Gly	Glu
		355					360					365			
Asp	Glu	Phe	Lys	Cys	Pro	lle	Lys	Glu	Glu	He	Ala	Leu	Thr	Ser	Gly
	370					375					380				
Glu	Trp	Glu	Val	Leu	Ala	Arg	His	Gly	Ser	Lys	Ile	Trp	Val	Asn	Glu
385					390					395					400
Glu	Thr	Lys	Leu	Val	Tyr	Phe	Gln	Gly	Thr	Lys	Asp	Thr	Pro	Leu	Glu
				405					410					415	
His	His	Leu	Tyr	Val	Val	Ser	Tyr	Glu	Ala	Ala	Gly	Glu	lle	Val	Arg
			420					425					430		
Leu	Thr	Thr	Pro	Gly	Phe	Ser	His	Ser	Cys	Ser	Met	Ser	Gln	Asn	Phe
		435					440					445			
Asp	Met	Phe	Val	Ser	His	Tyr	Ser	Ser	Val	Ser	Thr	Pro	Pro	Cys	Val
	450					455					460				
His	Val	Tyr	Lys	Leu	Ser	Gly	Pro	Asp	Asp	Asp	Pro	Leu	His	Lys	Gln
465					470					475					480
Pro	Arg	Phe	Trp	Ala	Ser	Met	Met	Glu		Ala	Ser	Cys	Pro	Pro	Asp
				485					490					495	
Tyr	Val	Pro		Glu	He	Phe	His		His	Thr	Arg	Ser		Val	Arg
			500					505					510		
Leu	Tyr	Gly	Met	He	Tyr	Lys		His	Ala	Leu	Gln		Gly	Lys	Lys
		515					520					525			
His		Thr	Val	Leu	Phe		Tyr	G] y	G] y	Pro		Val	Gln	Leu	Val
	530					535		_			540				
	Asn	Ser	Phe	Lys		lle	Lys	Tyr	Leu		Leu	Asn	Thr	Leu	
545					550					555					560

Ser Leu Gly Tyr Ala Val Val Ile Asp Gly Arg Gly Ser Cys Gln Arg Gly Leu Arg Phe Glu Gly Ala Leu Lys Asn Gln Met Gly Gln Val Glu Ile Glu Asp Gln Val Glu Gly Pro Gln Phe Val Ala Glu Lys Tyr Gly Phe Ile Asp Leu Ser Arg Val Ala Ile His Gly Trp Ser Tyr Gly Gly Phe Leu Ser Leu Met Gly Leu Ile His Lys Pro Gln Val Phe Lys Val Ala Ile Ala Gly Ala Pro Val Thr Val Trp Met Ala Tyr Asp Thr Gly Tyr Thr Glu Arg Tyr Met Asp Val Pro Glu Asn Asn Gln His Gly Tyr Glu Ala Gly Ser Val Ala Leu His Val Glu Lys Leu Pro Asn Glu Pro Asn Arg Leu Leu Ile Leu His Gly Phe Leu Asp Glu Asn Val His Phe Phe His Thr Asn Phe Leu Val Ser Gln Leu Ile Arg Ala Gly Lys Pro Tyr Gln Leu Gln 11e Tyr Pro Asn Glu Arg His Ser 11e Arg Cys Pro Glu Ser Gly Glu His Tyr Glu Val Thr Leu Leu His Phe Leu Gln Glu Tyr Leu <210> 4973

<211> 268

<212> PRT

<213> Homo sapiens

<400> 596

Met Ala Ala Ala Ile Thr Asp Met Ala Asp Leu Glu Glu Leu Ser Arg

Leu	Ser	Pro	Leu	Pro	Pro	Gly	Ser	Pro	Gly	Ser	Ala	Ala	Arg	Gly	Arg
			20					25					30		
Ala	Glu	Pro	Pro	Glu	Glu	Glu	Glu	Glu	Glu	Glu	Glu	Glu	Glu	Glu	Glu
		35					40					45			
Ala	Glu	Ala	Glu	Ala	Val	Ala	Ala	Leu	Leu	Leu	Asn	Gly	Gly	Ser	Gly
	50					55					60				
Gly	Gly	Gly	Gly	Gly	Gly	Gly	Gly	Gly	Val	Gly	Gly	Gly	Glu	Ala	Glu
65					70					75					80
Thr	Met	Ser	Glu	Pro	Ser	Pro	Glu	Ser	Ala	Ser	Gln	Ala	Gly	Glu	Asp
				85					90					95	
Glu	Asp	Glu	Glu	Glu	Asp	Asp	Glu	Glu	Glu	Glu	Asp	Glu	Ser	Ser	Ser
			100					105					110		
Ser	Gly	Gly	Gly	Glu	Glu	Glu	Ser	Ser	Ala	Glu	Ser	Leu	Val	Gly	Ser
		115					120					125			
Ser	Gly	Gly	Ser	Ser	Ser	Asp	Glu	Thr	Arg	Ser	Leu	Ser	Pro	Gly	Ala
	130					135					140				
Ala	Ser	Ser	Ser	Ser	Gly	Asp	G1 y	Asp	Gly	Lys	Glu	G1 y	Leu	Glu	Glu
145					150					155					160
Pro	Lys	Gly	Pro	Arg	Gly	Ser	Gln	Gly	Gly	Gly	Gly	Gly	Gly	Ser	Ser
				165					170					175	
Ser	Ser	Ser	Val	Val	Ser	Ser	Gly	Gly	Asp	Glu	Gly	Tyr	Gly	Thr	Gly
			180					185					190		
Gly	Gly	Gly	Ser	Ser	Ala	Thr	Ser	Gly	Gly	Arg	Arg	Gly	Ser	Leu	Glu
		195					200					205			
Met	Ser	Ser	Asp	Gly	Glu	Pro	Leu	Ser	Arg	Met	Asp	Ser	Glu	Asp	Ser
	210					215					220				
He	Ser	Ser	Thr	He	Met	Asp	Val	Asp	Ser	Thr	He	Ser	Ser	Gly	Arg
225					230					235					240
Ser	Thr	Pro	Ala	Met	Met	Asn	Gly	Gln	Gly	Ser	Thr	Thr	Ser	Ser	Ser
				245					250					255	
Lys	Asn	lle	Ala	Tyr	Asn	Cys	Cys	Trp	Asp	Gln	Cys				
			260					265							

<210> 4974 <211> 160 <212> PRT

<213> Homo sapiens

<400> 597

Met Leu Gly Glu Arg Pro Leu Cys Leu Thr Pro Gly Ala Lys Leu Gly
1 5 10 15

Leu His His Trp Ile Ile Arg Cys Phe Gln Arg Arg Pro Ser Pro His 20 25 30

Pro Asp Gln Gly Asn Trp Asp Asp Val Gly Pro Leu Thr Leu Ser Asp
35 40 45

Met Gln Pro Gly Trp Arg Leu Gly Gln Pro Pro Cys Arg Ser Glu Ser
50 55 60

Phe Ser Pro Cys His Leu Lys Ser Lys Ser Leu Leu Thr Gln Ala Leu 65 70 75 80

Pro Gly Met Cys Ser Pro Ala Thr Glu Pro Glu Ala Ala Leu Leu Leu 85 90 95

Ser Pro Val Gly Thr Ala Phe Gln Thr Gln Asn Thr Glu Phe Thr Ala 100 105 110

Ser Phe Gly Ile Phe Pro Asn Arg Leu Thr Val Pro Ile Ala Ser Asp 115 120 125

Pro Phe Gln Tyr Ser Cys Thr Arg Asn Val Met Val Lys Asn Lys Lys 130 135 140

Pro Pro Val Met Glu Gly Arg Asn Met Asn Gln Leu Lys Ala Phe His 145 150 155 160

<210> 4975

<211> 411

<212> PRT

<213> Homo sapiens

<400> 598

Met Thr Glu Met Ser Glu Lys Glu Asn Glu Pro Asp Asp Ala Ala Thr

1 5 10 15

His Ser Pro Pro Gly Thr Val Ser Ala Leu Gln Glu Thr Lys Leu Gln

20 25 30

Arg	Phe	Lys	Arg	Ser	Leu	Ser	Leu	Lys	Thr	Ile	Leu	Arg	Ser	Lys	Ser
		35					40					45			
Leu	Glu	Asn	Phe	Phe	Leu	Arg	Ser	Gly	Ser	Glu	Leu	Lys	Cys	Pro	Thr
	50					55					60				
Glu	Val	Leu	Leu	Thr	Pro	Pro	Thr	Pro	Leu	Pro	Pro	Pro	Ser	Pro	Pro
65					70					75					80
Pro	Thr	Ala	Ser	Asp	Arg	Gly	Leu	Ala	Thr	Pro	Ser	Pro	Ser	Pro	Cys
				85					90					95	
Pro	Val	Pro	Arg	Pro	Leu	Ala	Ala	Leu	Lys	Pro	Val	Arg	Leu	His	Ser
			100					105					110		
Phe	Gln	Glu	His	Val	Phe	Lys	Arg	Ala	Ser	Pro	Cys	Glu	Leu	Cys	His
		115					120					125			
Gln	Leu	He	Val	Gly	Asn	Ser	Lys	Gln	Gly	Leu	Arg	Cys	Lys	Met	Cys
	130					135					140				
Lys	Val	Ser	Val	His	Leu	Trp	Cys	Ser	Glu	Glu	lle	Ser	His	Gln	Gln
145					150					155					160
Cys	Pro	Gly	Lys	Thr	Ser	Thr	Ser	Phe	Arg	Arg	Asn	Phe	Ser	Ser	Pro
				165					170					175	
Leu	Leu	Val	His	Glu	Pro	Pro	Pro	Val	Cys	Ala	Thr	Ser	Lys	Glu	Ser
			180					185					190		
Pro	Pro	Thr	Gly	Asp	Ser	Gly	Lys	Val	Asp	Pro	Val	Tyr	Glu	Thr	Leu
		195					200					205			
Arg	Tyr	Gly	Thr	Ser	Leu	Ala	Leu	Met	Asn	Arg	Ser	Ser	Phe	Ser	Ser
	210					215					220				
Thr	Ser	Glu	Ser	Pro	Thr	Arg	Ser	Leu	Ser	Glu	Arg	Asp	Glu	Leu	Thr
225					230					235					240
Glu	Asp	Gly	Glu	Gly	Ser	lle	Arg	Ser	Ser	Glu	Glu	Gly	Pro	Gly	Asp
				245					250					255	
Ser	Ala	Ser	Pro	Val	Phe	Thr	Ala	Pro	Ala	Glu	Ser	Glu	Gly	Pro	Gly
			260					265					270		
Pro	Glu	Glu	Lys	Ser	Pro	G]y	Gln	Gln	Leu	Pro	Lys	Ala	Thr	Leu	Arg
		275					280					285			
Lys	Asp	Val	Gly	Pro	Met	Tyr	Ser	Tyr	Val	Ala	Leu	Tyr	Lys	Phe	Leu
	290					295					300				
Pro	G1n	Glu	Asn	Asn	Asp	Leu	Ala	Leu	Gln	Pro	Gly	Asp	۸rg	lle	Met
305					310					315					320

Leu Val Asp Asp Ser Asn Glu Asp Trp Trp Lys Gly Lys Ile Gly Asp Arg Val Gly Phe Phe Pro Ala Asn Phe Val Gln Arg Val Arg Pro Gly Glu Asn Val Trp Arg Cys Cys Gln Pro Phe Ser Gly Asn Lys Glu Gln Gly Tyr Met Ser Leu Lys Glu Asn Gln Ile Cys Val Gly Val Gly Arg Ser Lys Asp Ala Asp Gly Phe Ile Arg Val Ser Ser Gly Lys Lys Arg Gly Leu Val Pro Val Asp Ala Leu Thr Glu Ile

<210> 4976

<211> 129

<212> PRT

<213> Homo sapiens

<400> 599

Met Leu Ser Ala Gly Gly Cys Ile Leu Leu Pro Val His Ile Val Cys Ala Trp Asp Gly His Glu Ser Gly Gly Arg Ala Gly Val Leu Ala Leu Gly Tyr Phe Gly Glu Thr 11e Arg Phe Lys Glu Gly Asn Thr Phe Ser Gly Pro Val Pro Gln Asn Thr Thr lle Thr Arg Gly Arg His Leu Ala Glu Leu Ser Pro Thr Gly Thr Ser Gly Pro Trp Phe Leu His Pro Gln Ser Ala Ser Phe His Ser Arg Gln Asn Gln Gly Ile Phe Gln Gly Gln Val Ser His Pro Leu Gly Ala Gly Ile Leu Ser His Ser Lys Thr Asn Phe Gln Val Pro Gly Leu Leu Gly Asn Pro Lys Arg Leu Ser Pro Ala

Glu

<210> 4977

```
<211> 156
<212> PRT
<213> Homo sapiens
<400> 600
Met Gln Thr Glu Asp Ile Arg Leu Glu Pro Asp Leu Tyr Glu Ala Cys
Lys Ser Asp Ile Lys Asn Phe Cys Ser Ala Val Gln Tyr Gly Asn Ala
             20
                                 25
                                                      30
Gln lle Ile Glu Cys Leu Lys Glu Asn Lys Lys Gln Leu Ser Thr Arg
                             40
                                                  45
Cys His Gln Lys Val Phe Lys Leu Gln Glu Thr Glu Met Met Asp Pro
                         55
                                              60
Glu Leu Asp Tyr Thr Leu Met Arg Val Cys Lys Gln Met Ile Lys Arg
                     70
 65
                                          75
                                                              80
Phe Cys Pro Glu Ala Asp Ser Lys Thr Met Leu Gln Cys Leu Lys Gln
                                      90
Asn Lys Asn Ser Glu Leu Met Asp Pro Lys Cys Lys Gln Met Ile Thr
            100
                                 105
                                                     110
Lys Arg Gln Ile Thr Gln Asn Thr Gly Lys Ile Leu Ala Trp Leu Ser
        115
                            120
                                                 125
Trp Pro Arg Gly Val Ser Glu Lys Glu Phe Ser Gly Cys Arg Val Thr
                        135
Cys Ser Asn Ser Gln Gly Phe Val Ala Trp Glu Phe
145
                    150
                                         155
```

<210> 4978

<211> 183

<212> PRT

<213> Homo sapiens

Met Glu Leu Ala Ala Leu Gly Leu Ser Pro Cys Pro Arg Leu Leu His Ala Glu Leu Leu Pro Gly Leu Leu Thr Val Phe Ser Leu Arg Phe Leu Gln Asp Tyr Gly Gly Tyr Leu Ser Thr Tyr Ile Leu Pro Ala Lys Gly Glu Asn Gln Gly Gln Thr Phe Thr Cys Gly Ser Ala Leu Ser Pro Ile Thr Asp Phe Lys Leu Tyr Ala Ser Ala Phe Ser Glu Arg Tyr Leu Gly Leu His Gly Thr Arg Asp Asn Ala Val Pro Ala Ala Ala Pro Pro Pro Gly Ala Arg Glu Thr Gly Thr Pro Arg Pro Leu Pro Gln Gly Thr Glu Gln Arg Met Val Ala Ala Gly Pro Thr Arg Ala His Arg Thr Pro Ala Pro Arg Phe Gln Pro Pro Ser Gly Ser Met Arg Pro Ala His Thr Ser Leu Cys Val Pro Val Arg Asp Ile Thr Pro Cys Leu Thr Ser Gln Cys His Gly Arg Ser Ser Tyr Ser Thr lle Val Leu Ala Val Arg Val His lle Trp Ala Cys Tyr Phe Leu <210> 4979 <211> 579 <212> PRT <213> Homo sapiens

Met Lys Asp Lys Ser Asn Gly Leu Glu Ser Gln Val Asn Gln Cys Asp

<400> 601

<400> 602

Lys	Met	Leu	Gly	Gly	Asp	Ala	Leu	Val	Thr	Asp	Leu	Leu	Val	Asp	Phe
			20					25					30		
Cys	Gly	Ser	Arg	Ser	Gly	Val	Glu	Ile	Pro	Arg	Thr	Pro	Gln	Leu	Tyr
		35					40					45			
Val	Ala	His	Glu	He	Gly	Thr	He	Lys	Thr	Val	Thr	Pro	Pro	Glu	Asp
	50					55					60				
Arg	Asp	Ser	Glu	Ser	Gly	Val	Val	Gly	Gly	Gln	Gly	Thr	Leu	Gln	Glu
65					70					75					80
Pro	Gly	Phe	Gly	Glu	Ala	Ser	Glu	Ala	Ile	Ser	Val	Ser	Arg	Asn	Arg
				85					90					95	
Gln	Pro	lle	Pro	Leu	Leu	Met	Asn	Lys	Glu	Asn	Ser	Thr	Lys	Thr	Ser
			100					105					110		
Lys	Val	Glu	Leu	Thr	Leu	Ala	Ser	Pro	Tyr	Met	Lys	Gln	Glu	Lys	Glu
		115					120					125			
Glu	Glu	Lys	Glu	Gly	Phe	Ser	Glu	Ser	Asp	Phe	Ser	Asp	Gly	Asn.	Thr
	130					135					140				
Ser	Ser	Asn	Ala	Glu	Ser	Trp	Arg	Asn	Pro	Ser	Ser	Ser	Glu	Glu	Glu
145					150					155					160
Pro	Ser	Pro	Val	Leu	Lys	Thr	Leu	Glu	Arg	Ser	Ala	Ala	Arg	Lys	Met
				165					170					175	
Pro	Ser	Lys	Ser	Leu	Glu	Asp	He	Ser	Ser	Asp	Ser	Ser	Asn	Gln	Ala
			180					185					190		
Lys	Val	Asp	Asn	Gln	Pro	Glu	G] u	Leu	Val	Arg	Ser	Ala	Glu	Asp	Asp
		195					200					205			
G]u	Lys	Pro	Asp	Gln	Lys	Pro	Val	Thr	Asn	Glu	Cys	Val	Pro	Arg	He
	210					215					220				
Ser	Thr	Val	Pro	Thr	Gln	Pro	Asp	Asn	Pro	Phe	Ser	His	Pro	Asp	Lys
225					230					235					240
Leu	Lys	Arg	Met	Ser	Lys	Ser	Val	Pro	Ala	Phe	Leu	Gln	Asp	Glu	Val
				245					250					255	
Ser	Gly	Ser	Val	Met	Ser	Va]	Tyr	Ser	Gly	Asp	Phe	G1 y	Asn	Leu	Glu
•			260					265					270		
Val	Lys	Gly	Asn	He	Gln	Phe		He	Glu	Tyr	Val	Glu	Ser	Leu	Lys
		275					280					285			
Glu		His	Va]	Phe	Val	Ala	Gln	Cys	Lys	Asp	Leu	Ala	Ala	Val	Asp
	290					295					300				

Val	Lys	Lys	Gln	Arg	Ser	Asp	Pro	Tyr	Val	Lys	Ala	Tyr	Leu	Leu	Pro
305					310					315					320
Asp	Lys	Gly	Lys	Met	Gly	Lys	Lys	Lys	Thr	Leu	Val	Val	Lys	Lys	Thr
				325					330					335	
Leu	Asn	Pro	Val	Tyr	Asn	Glu	lle	Leu	Arg	Ty.r	Lys	He	Glu	Lys	Gln
			340					345					350		
He	Leu	Lys	Thr	Gln	Lys	Leu	Asn	Leu	Ser	He	Trp	His	Arg	Asp	Thr
		355					360					365			
Phe	Lys	Arg	Asn	Ser	Phe	Leu	G] y	Glu	Val	${\tt Glu}$	Leu	Asp	Leu	Glu	Thr
	370					375					380				
Trp	Asp	Trp	Asp	Asn	Lys	Gln	Asn	Lys	Gln	Leu	۸rg	Trp	Tyr	Pro	Leu
385					390					395					400
Lys	Arg	Lys	Thr	Ala	Pro	Val	Ala	Leu	Glu	Ala	Glu	Asn	Arg	Gly	Glu
				405					410				•	415	
Met	Lys	Leu	Ala	Leu	Gln	Tyr	Val	Pro	Glu	Pro	Val	Pro	Gly	Lys	Lys
			420					425					430		
Leu	Pro	Thr	Thr	Gly	Glu	Val	His	lle	Trp	Val	Lys	Glu	Cys	Leu	Asp
		435					440					445			
Leu	Pro	Leu	Leu	Arg	Gly	Ser	His	Leu	Asn	Ser	Phe	Val	Lys	Cys	Thr
	450					455					460				
He	Leu	Pro	Asp	Thr	Ser	Arg	Lys	Ser	Arg	Gln	Lys	Thr	Arg	Ala	Val
465					470					475					480
G] y	Lys	Thr	Thr	Asn	Pro	lle	Phe	Asn	His	Thr	Met	Val	Tyr	Asp	Gly
				485					490					495	
Phe	Arg	Pro	Glu	Asp	Leu	Met	Glu	Ala	Cys	Val	Glu	Leu	Thr	Va]	Trp
			500					505					510		
Asp	llis		Lys	Leu	Thr	Asn		Phe	Leu	Gly	Gly		Arg	lle	Gly
		515					520					525			
Phe		Thr	G1 y	Lys	Ser		G1 y	Thr	Glu	Val		Trp	Met	Asp	Ser
	530					535					540				
	Ser	Glu	Glu	Val	Ala	Leu	Trp	Glu	Lys		Val	Asn	Ser	Pro	
545		_			550					555					560
Thr	Trp	He	Glu		Thr	Leu	Pro	Leu		Met	Leu	Leu	He		Lys
	_			565					570					575	
110	Sar	Lvc													

```
<211> 261
<212> PRT
<213> Homo sapiens
<400> 603
Met Arg Ser Leu Leu Leu Ser Ala Phe Cys Leu Leu Glu Ala Ala
                                                          15
 1
                  5
                                     10
Leu Ala Ala Glu Val Lys Lys Pro Ala Ala Ala Ala Ala Pro Gly Thr
                                 25
Ala Glu Lys Leu Ser Pro Lys Ala Ala Thr Leu Ala Glu Arg Ser Ala
                             40
Gly Leu Ala Phe Ser Leu Tyr Gln Ala Met Ala Lys Asp Gln Ala Val
                         55
                                             60
Glu Asn Ile Leu Val Ser Pro Val Val Val Ala Ser Ser Leu Gly Leu
                     70
                                         75
Val Ser Leu Gly Gly Lys Ala Thr Thr Ala Ser Gln Ala Lys Ala Val
                 85
                                     90
                                                          95
Leu Ser Ala Glu Gln Leu Arg Asp Glu Glu Val His Ala Gly Leu Gly
                                105
Glu Leu Leu Arg Ser Leu Ser Asn Ser Thr Ala Arg Asn Val Thr Trp
        115
                            120
                                                 125
Lys Leu Gly Ser Arg Leu Tyr Gly Pro Ser Ser Val Ser Phe Ala Asp
    130
                        135
                                            140
Asp Phe Val Arg Ser Ser Lys Gln His Tyr Asn Cys Glu His Ser Lys
                    150
                                       155
lle Asn Phe Arg Asp Lys Arg Ser Ala Leu Gln Ser lle Asn Glu Trp
                165
                                    170
                                                         175
Ala Ala Gln Thr Thr Asp Gly Lys Leu Pro Glu Val Thr Lys Asp Val
                                185
Glu Arg Thr Asp Gly Ala Leu Leu Val Asn Ala Met Phe Phe Lys Arg
        195
                            200
                                                205
Glu Ser Gly Ala Arg Ser Gly Val Leu Leu Leu Pro Gly Pro Pro
    210
                        215
                                            220
```

<210> 4980

Ala Arg Val Arg Thr Thr Phe Arg Ala Leu His Ser Ser Leu Pro Leu lle Tyr Ala Val Thr Thr Gln Gly Gly Arg Thr Val Thr Gln Leu Phe Val Gln Thr Gly Asn

<210> 4981 <211> 281 <212> PRT <213> Homo sapiens

<400> 604

Met Ala Glu Ala Ala Pro Ala Arg Asp Pro Glu Thr Asp Lys His Thr Glu Asp Gln Ser Pro Ser Thr Pro Leu Pro Gln Pro Ala Ala Glu Lys Asn Ser Tyr Leu Tyr Ser Thr Glu IIe Thr Leu Trp Thr Val Val Ala Ala Ile Gln Ala Leu Glu Lys Lys Val Asp Ser Cys Leu Thr Arg Leu Leu Thr Leu Glu Gly Arg Thr Gly Thr Ala Glu Lys Lys Leu Ala Asp Cys Glu Lys Thr Ala Val Glu Phe Gly Asn Gln Leu Glu Gly Lys Trp Ala Val Leu Gly Thr Leu Leu Gln Glu Tyr Gly Leu Leu Gln Arg Arg Leu Glu Asn Val Glu Asn Leu Leu Arg Asn Arg Asn Phe Trp lle Leu Arg Leu Pro Pro Gly Ser Lys Gly Glu Ala Pro Lys Val Pro Val Thr Phe Asp Asp Val Ala Val Tyr Phe Ser Glu Leu Glu Trp Gly Lys Leu Glu Asp Trp Gln Lys Glu Leu Tyr Lys His Val Met Arg Gly Asn Tyr

Glu Thr Leu Val Ser Leu Asp Tyr Ala Ile Ser Lys Pro Asp Ile Leu 180 185 Thr Arg Ile Glu Arg Gly Glu Glu Pro Cys Leu Asp Arg Trp Gly Gln 200 205 195 Glu Lys Gly Asn Glu Val Glu Val Gly Arg Pro Arg Met Met Gly Thr 215 220 Gly Leu Pro Pro Tyr Pro Glu His Leu Thr Ser Pro Leu Ser Pro Ala 230 235 240 Gln Glu Glu Leu Lys Glu Gly Gln Ala Pro Lys Gln Gln Gln Asp Ser 250 245 255 Glu Ala Arg Val Ala Pro Ala Gly Pro Glu Ala Gly Leu Ala Leu Arg 265 Thr Asp Leu Gln Gly Glu Ala Gln lle 275 280

<210> 4982

<211> 126

<212> PRT

<213> Homo sapiens

<400> 605

 Met
 Ser
 Pro
 Pro
 Leu
 Leu
 Pro
 Pro
 Gly
 Pro
 Ser
 His
 His
 His
 Val

 1
 5
 10
 10
 15
 15
 15

 Ser
 Pro
 Gly
 Leu
 His
 Ala
 Pro
 Pro
 Trp
 Ser
 Pro
 His
 Phe
 Cys

 Pro
 Gly
 Arg
 Gly
 Trp
 Cys
 Val
 Leu
 His
 Pro
 Ala
 Thr
 Gln
 Gln
 Ser
 Glu

 Gly
 Ser
 Ser
 Tyr
 Asn
 Ile
 Ser
 Gln
 Ile
 Thr
 Ser
 Leu
 Phe
 Ala
 Gln
 Asn

 50
 55
 55
 60
 60
 45
 45
 45

Pro Ala Met Ala Pro 11e Ser Phe Arg Val Asn Ser Lys Ala Pro Tyr 65 70 75 80

Leu Pro Thr Ala Pro His Ser Arg Thr Phe Pro Pro Val Val Leu Pro 85 90 95

Leu Val Pro Ser Ile Pro Ala Thr Gly Ile Ser Leu Leu Ala Pro Glu 100 105 110 Leu Thr Arg Tyr Gly Ala Ala Ser Gly Pro Leu His Leu Gln 115 120 125

<210> 4983

<211> 266

<212> PRT

<213> Homo sapiens

<400> 606

Met Val Met Ala Asp Gln Asn Gln Val Trp Val Gly Ser Glu Asp Ser

1 5 10 15

Val Ile Tyr 11e Ile Asn Val His Ser Met Ser Cys Asn Lys Gln Leu 20 25 30

Thr Ala His Cys Ser Ser Val Thr Asp Leu Ile Val Gln Asp Gly Gln
35 40 45

Glu Ala Pro Ser Asn Val Tyr Ser Cys Ser Met Asp Gly Met Val Leu
50 55 60

Val Trp Asn Val Ser Thr Leu Gln Val Thr Ser Arg Phe Gln Leu Pro
65 70 75 80

Arg Gly Gly Leu Thr Ser Ile Arg Leu His Gly Gly Arg Leu Trp Cys
85 90 95

Cys Thr Gly Asn Ser Ile Met Val Met Lys Met Asn Gly Ser Leu His 100 105 110

Gln Glu Leu Lys 11e Glu Glu Asn Phe Lys Asp Thr Ser Thr Ser Phe 115 120 125

Leu Ala Phe Gln Leu Leu Pro Glu Glu Glu Gln Leu Trp Ala Ala Cys 130 135 140

Ala Gly Arg Ser Glu Val Tyr Ile Trp Ser Leu Lys Asp Leu Ala Gln 145 150 155 160

Pro Pro Gln Arg Val Pro Leu Glu Asp Cys Ser Glu lle Asn Cys Met
165 170 175

lle Arg Val Lys Lys Gln Val Gly Trp Arg Ala Arg His Pro Gln His 180 185 190

Pro Arg Gln Val Ser Leu Ala Leu Ala Ala Ser Pro Cys Ser Arg Glu 195 200 205 Pro Ala Ala Arg Pro Arg Ala Leu Leu Pro Ser Pro Leu Arg Val Pro Leu Leu Thr Gly Thr Cys Arg Val Gly Ala Asn Gly Pro Thr Gly His His Val Leu Val Ser Trp Cys Gly Pro Ser Thr Pro Gln Arg Ile Pro Pro Asn Trp Met Ala Ser Asn Asp Arg Thr

<210> 4984

<211> 251

<212> PRT

<213> Homo sapiens

<400> 607

Met Gly Met Gly Thr Leu Ala Trp Gly Gln Pro Leu Leu Pro Arg Cys Leu Cys Pro Arg Ala Gly Gln Arg Gln Pro Val Val Thr Ala Ala Val Ala Ala Pro Gln Leu Thr Met Asn Asp Phe Ser Val His Arg Ile Ile Gly Arg Gly Gly Phe Gly Glu Val Tyr Gly Cys Arg Lys Ala Asp Thr Gly Lys Met Tyr Ala Met Lys Cys Leu Asp Lys Lys Arg Ile Lys Met Lys Gln Gly Glu Thr Leu Ala Leu Asn Glu Arg lle Met Leu Ser Leu Val Ser Thr Gly Asp Cys Pro Phe 11e Val Cys Met Ser Tyr Ala Phe His Thr Pro Asp Lys Leu Ser Phe Ile Leu Asp Leu Met Asn Gly Gly Asp Leu His Tyr His Leu Ser Gln His Gly Val Phe Ser Glu Ala Asp

Met Arg Phe Tyr Ala Ala Glu IIe IIe Leu Gly Leu Glu His Met His

Asn Arg Phe Val Val Tyr Arg Asp Leu Lys Val Ser Ala Pro Ala Val Pro Arg Leu Asp Leu Arg Gly Cys Pro Leu Leu Pro Leu Asp Ile Pro Ala Thr Arg Pro Arg Gly Val Gly Leu Leu Gly His Gly Arg Pro Val Ser Ser His Leu Arg Pro Cys Pro Ser His Arg Ala Thr Leu Trp Val Gln Val Val Ala Gly Asp Arg Arg Glu Asp Pro His Leu Cys Pro Phe Phe Gly Tyr Pro Ser Ser Ser Pro Val Lys Gln

<210> 4985

<211> 350

<212> PRT

<213> Homo sapiens

<400> 608

Met Met Glu Ser Ser Glu Leu Thr Pro Lys Gln Glu 11e Phe Lys Gly Ser Glu Ser Ser Asn Ser Thr Ser Gly Gly Leu Phe Gly Val Val Pro Gly Gly Thr Glu Thr Gly Asp Val Cys Glu Asp Thr Phe Lys Glu Leu Glu Gly Gln Pro Ser Asn Glu Glu Gly Ser Arg Leu Glu Ser Asp Phe Leu Glu 11e 11e Asp Glu Asp Lys Lys Lys Ser Thr Lys Asp Arg Tyr Glu Glu Tyr Lys Glu Val Glu Glu His Pro Pro Leu Ser Ser Pro Val Glu His Glu Gly Val Leu Lys Gly Gln Lys Ser Tyr Arg Cys Asp Glu Cys Gly Lys Ala Phe Tyr Trp Ser Ser His Leu Ile Gly His Arg

Arg	He	His	Thr	Gly	Glu	Lys	Pro	Tyr	Glu	Cys	Asn	Glu	Cys	Gly	Lys
	130					135					140				
Thr	Phe	Λrg	G1n	Thr	Ser	Gln	Leu	He	Val	His	Leu	Arg	Thr	His	Thr
145					150					155					160
G1 y	Glu	Lys	Pro	Tyr	Glu	Cys	Ser	Glu	Cys	Gly	Lys	Лlа	Tyr	Arg	His
				165					170					175	
Ser	Ser	His	Leu	He	Gln	His	Gln	۸rg	Leu	His	Asn	Gly	Glu	Lys	Pro
			180					185					190		
Tyr	Lys	Cys	Asn	Glu	Cys	Ala	Lys	Ala	Phe	Asn	Gln	Ser	Ser	Lys	Leu
		195					200					205			
Phe	Asp	His	Gln	Arg	Thr	His	Thr	Gly	Glu	Lys	Pro	Tyr	Glu	Cys	Lys
	210					215					220				
Glu	Cys	Gly	Ala	Ala	Phe	Ser	Arg	Ser	Lys	Asn	Leu	Val	Arg	His	Gln
225					230					235					240
Phe	Leu	His	Thr	Gly	Lys	Lys	Pro	Tyr	Lys	Cys	Asn	Glu	Cys	Gly	Arg
				245					250					255	
Ala	Phe	Cys	Ser	Asn	Arg	Asn	Leu	Ile	Asp	His	Gln	Arg	Thr	His	Thr
			260					265					270		
Gly	Glu	Lys	Pro	Tyr	Lys	Cys	Asn	Glu	Cys	Gly	Lys	Ala	Phe	Ser	Arg
		275					280					285			
Ser	Lys	Cys	Leu	lle	Arg	His	Gln	Ser	Leu	His	Thr	Gly	Glu	Lys	Pro
	290					295					300				
Tyr	Lys	Cys	Ser	Glu	Cys	Gly	Lys	Ala	Phe	Asn	Gln	He	Ser	Gln	Leu
305					310					315					320
Val	Glu	His	Glu	Arg	lle	His	Thr	G1 y	Glu	Lys	Pro	Phe	Lys	Cys	Ser
				325					330					335	
Glu	Cys	Gly	Lys	Ala	Phe	Gly	Leu	Ser	Lys	Cys	Leu	He	Arg		
			340					345					350		

<211> 191

<212> PRT

<213> Homo sapiens

Met Tyr Ala Lys Arg Pro Glu Ser Phe Ser Cys Lys Leu Ala Met

1 10 15 Ser Lys Lys Val Thr Pro Ala Ser Thr Gln Cys Val Arg Phe Pro Phe 20 25 Pro Ser Thr Val Val Cys Lys Lys Asn Leu Asp Ser Thr Thr Val Ala 40 45 Val His Gly Glu Glu Ile Tyr Cys Lys Ser Cys Tyr Gly Lys Lys Tyr 50 55 60 Gly Pro Lys Gly Tyr Gly Tyr Gly Gln Gly Ala Gly Thr Leu Ser Thr 75 Asp Lys Gly Glu Ser Leu Gly lle Lys His Glu Glu Ala Pro Gly His Arg Pro Thr Thr Asn Pro Asn Ala Ser Lys Phe Ala Gln Lys Ile Glv 100 105 110 Gly Ser Glu Arg Cys Pro Arg Cys Ser Gln Ala Val Tyr Ala Ala Glu 120 Lys Val Ile Gly Ala Gly Lys Ser Trp His Lys Ala Cys Phe Arg Cys 130 135 140 Ala Lys Cys Gly Lys Gly Leu Glu Ser Thr Thr Leu Ala Asp Lys Asp 150 155 160 Gly Glu Ile Tyr Cys Lys Gly Cys Tyr Ala Lys Asn Phe Gly Pro Lys 170 Gly Phe Gly Phe Gly Gln Gly Ala Gly Ala Leu Val His Ser Glu 180 185 190

<210> 4987

<211> 482

<212> PRT

<213> Homo sapiens

<400> 610

Met Trp Thr Val Pro Ser Phe Thr Asn Asp Ser Tyr Gln Val Tyr Asn

1 5 10 15

Val Phe Ser Thr Asn Ser Phe Gln Leu Leu Thr Val Lys Arg Thr Pro

			20					25					30		
His	Glu	Λla	Trp	Arg	Val	Pro	Leu	Thr	Thr	Lys	Thr	Asn	Lys	Thr	Lys
		35					40					45			
Gly	Leu	Pro	Asp	Cys	Pro	Lys	Lys	Pro	Thr	Asn	Gly	Pro	Phe	lle	Va]
	50					55					60				
Thr	Ser	He	Leu	Trp	Asp	Asn	Cys	Asn	Ala	Pro	Lys	Ala	Val	Val	Leu
65					70					75					80
Gln	Thr	Leu	Ala	Met	Gly	lle	Val	11e	Asp	Trp	Ala	Pro	Lys	Gly	His
				85					90					95	
Tyr	Trp	Gln	Asp	Cys	Ser	Ser	Lys	Asn	Thr	Leu	Cys	Ser	Glu	Phe	Ile
			100					105					110		
Tyr	Ser	Leu	Asp	Tyr	He	Glu	His	G1 y	Trp	Gln	Ser	Tyr	Thr	Met	۸rg
		115					120					125			
Gln	Arg	Val	Ser	Pro	Tyr	Pro	Phe	Lys	Trp	Met	Asp	Thr	Gly	lle	Ala
	130					135					140				
Pro	Pro	Arg	Pro	Lys	He	He	His	Pro	Phe	Phe	Thr	Pro	Glu	His	Pro
145					150					155					160
Glu	Leu	Trp	Lys	Leu	Ala	Ala	Ala	Leu	Ser	Gly	lle	Lys	Ile	Trp	Asn
				165					170					175	
Thr	Thr	Tyr	Gln	Leu	Leu	Arg	Thr	Lys	Thr	Lys	Thr	Pro	Thr	Phe	Asn
			180					185					190		
Hle	Thr		He	Ser	Glu	Trp	Val	lle	Pro	He	Arg	Ser	Cys	Val	Lys
		195					200					205			
Pro		Tyr	Met	Leu	Leu	Val	G] y	Asn	He	He		Met	Pro	Asp	Ala
0.1	210			_		215					220				
	Thr	He	Glu	Cys		Asn	Cys	Lys	Leu		Thr	Cys	He	Asp	
225	Di		n	m.	230					235					240
Inr	Phe	Asn	Pro		Thr	Ser	He	Leu		Val	Arg	Ala	Arg		Gly
W - 3	т	11	D	245	C	,			250 D		0.1			255	
vai	irp	116		vai	Ser	Leu	HIS		Pro	lrp	Glu	Ser		Pro	Ser
T1a	u; a	11.	260 V-1	۸	C1	W = 1	1	265		11			270	TI.	
116	nis		vai	ASI	GIU	Val		Lys	Asp	116	Leu		Arg	ınr	Lys
Arc	Pho	275	Db.	Th	1	11.	280	Va 1	Lace	д1 -	C1	285	1	λ1.	V = 1
ni g	290	116	тпе	HII	reu	11e 295	V19	val	Leu	ита		Leu	ren	ита	118 v
Thr		Thr	Δla	Ala	Thr	Ala	C1v	Val	Alo	116	300	Sor	Son	Vol	Clr

305					310					315					320
Thr	Ala	His	Tyr	Val	Glu	Λla	Cys	Gln	Lys	Asn	Ser	Ser	Arg	Leu	Trp
				325					330					335	
Asn	Ser	Gln	Ala	Gln	Ile	Asp	Gln	Lys	Leu	Ala	Asn	Gln	Пe	Asn	Asp
			340					345					350		
Leu	Arg	G1n	Ser	Val	Thr	Trp	Leu	G1 y	Asp	Arg	Val	Met	Asn	Leu	Gln
		355					360					365			
His	Arg	Met	Gln	Leu	Gln	Cys	Asp	Trp	Asn	Thr	Ser	Asp	Tyr	Cys	He
	370					375					380				
Thr	Pro	Tyr	Ala	Tyr	Asn	Gln	Asp	Gln	His	Ser	Trp	Glu	Asn	Val	Ser
385					390					395					400
Arg	His	Leu	Lys	Ala	Trp	Asp	Asp	Asn	Leu	Thr	Leu	Asp	He	Ser	Gln
				405					410					415	
Leu	Lys	Glu	Gln	lle	Phe	Glu	Ala	Ser	Gln	Val	His	Leu	Ser	Thr	Val
			420					425					430		
Pro	Gly	Ser	His	Ile	Phe	Glu	Gly	He	Thr	Lys	Gln	Leu	Pro	Asp	Phe
		435					440					445			
Asn	Pro	Phe	Lys	Trp	Leu	Lys	Pro	Val	Arg	Gly	Ser	Leu	Leu	Leu	Leu
	450					455					460				
Ala	Leu	Leu	He	Leu	Val	Cys	Leu	Cys	Cys	Leu	Leu	Leu	Val	Cys	Arg
465					470					475					480
Cys	Leu														

<211> 465

<212> PRT

<213> Homo sapiens

<400> 611

		35					40					45			
Pro	Cys	Pro	Arg	Thr	Val	Pro	Ser	Ala	Ser	Gly	Ala	Trp	Pro	Pro	Cys
	50					55	`				60				
Asn	Ser	Ser	Gly	Ala	Val	Arg	Ser	Phe	Ser	Leu	Glu	Ala	Ala	Glu	Va]
65					70					75					80
Leu	Val	Glu	His	Thr	Cys	His	Arg	Gly	Λla	Val	Thr	Gly	Leu	Thr	Ala
				85					90					95	
Thr	Pro	Asp	Gly	Arg	Leu	Leu	Phe	Ser	Ser	Cys	Ser	Gln	Gly	Ser	Leu
			100					105					110		
Ala	Gln	Tyr	Ser	Cys	Ala	Asp	Pro	Gln	Trp	His	Val	Leu	Arg	Val	Ala
		115					120					125			
Ala	Asp	Met	Va]	Cys	Pro	Asp	Ala	Pro	Ala	Ser	Pro	Ser	Ala	Leu	Ala
	130					135					140				
Val	Ser	Arg	Asp	Gly	Arg	Leu	Leu	Ala	Phe	Val	Gly	Pro	Ser	Arg	Cys
145					150					155					160
Thr	Val	Thr	Val	Met	Gly	Ser	Ala	Ser	Leu	Asp	Glu	Leu	Leu	Arg	Val
				165					170					175	
Asp	lle	Gly		Leu	Asp	Leu	Ala		Ser	Arg	Leu	Asp		Ala	Met
		_	180		_	_		185					190		
Ala	Val		Phe	Gly	Pro	Ala		Leu	Gly	Gln	Leu		Val	Ser	Thr
<u> </u>	0	195				., 1	200					205			
Ser		Asn	Arg	Val	Val		Leu	Asp	Ala	Val		Gly	Arg	Пе	He
A	210		D.	C1	V 1	215	D	C1	D	C	220 D	C		TI	
	GIU	Leu	Pro	G1 y		HIS	Pro	GIU	Pro		Pro	Ser	Leu	Thr	
225 San	C1	Aan	A10	A 22.00	230 Dho	Lau	Lau	11.	A1.	235	C1	۸	TL	11.	240
sei	Glu	nsp	мта	245	rne	Leu	Leu	116	250	Ala	GIY	Arg	inr	He	Lys
Val	Trn	Acn	Tyr		The	Cln	Λlα	San		C1	Dno	Cln.	Val	255 Turn	II.
vai	пр	nsp	260	мта	1111	OTH	мла	265	F10	GIŸ	110	GIN	270	Tyr	116
Glv	Hic	Ser		Pro	Val	Gln	Ala		Ala	Pho	Sor	Pro		Gln	Gln
013	1113	275	Olu	1,0	, 41	OIII	280	101	та	THE	361	285	nsp	0111	OIII
Gln	Val		Ser	Ala	G1v	Asp		Val	Phe	l eu	Tro		Val	Leu	Ala
••	290	200	~ ~ .		<i>-1</i> ,	295				Dea	300	9	. (1,1	200	. 3 2 CI
Thr		Glu	Ser	Asp	Gln		Phe	Pro	G] v	Ala		Pro	Ala	Cys	Lvs
305					310				,	315				- , -	320

```
Thr Gly Pro Gly Ala Gly Pro Leu Glu Asp Ala Ala Ser Arg Ala Ser
                                    330
                325
Glu Leu Pro Arg Gln Gln Val Pro Lys Pro Cys Gln Ala Ser Pro Pro
            340
                                                    350
                                345
Arg Leu Gly Val Cys Ala Arg Pro Pro Glu Gly Gly Asp Gly Ala Arg
                            360
                                                365
Asp Thr Arg Asn Ser Gly Ala Pro Arg Thr Thr Tyr Leu Ala Ser Cys
                        375
Lys Ala Phe Thr Leu Ala Arg Val Ser Cys Ser Pro His Ser Ala Lys
                    390
                                        395
385
Gly Thr Cys Pro Pro Pro Ala Ser Gly Gly Trp Leu Arg Leu Lys Ala
                405
                                    410
Val Val Gly Tyr Ser Gly Asn Gly Arg Ala Asn Met Val Trp Arg Pro
            420
                                425
                                                    430
Asp Thr Gly Gly Gly Gln Glu Pro Thr Pro Thr Pro Ser Gln Asp Ala
                            440
                                                445
Ala Ala Arg Gly Pro Ala Val Ser Thr Pro Arg Pro Gly Pro Gly Gly
                        455
                                            460
Lys
465
<210> 4989
<211> 134
<212> PRT
<213> Homo sapiens
<400> 612
Met Gly Lys Lys Gln Ser Arg Lys Thr Gly Asn Ser Lys Lys Gln Ser
 1
                                     10
Ala Ser Pro Pro Pro Lys Glu Cys Ser Ser Ser Pro Ala Thr Glu Gln
                                 25
Ser Trp Met Glu Asn Asp Phe Asp Glu Leu Arg Glu Ala Gly Phe Arg
         35
                                                 45
Arg Ser Asn Tyr Ser Glu Leu Arg Glu Asp Ile Gln Thr Lys Gly Lys
```

Lys Val Glu Asn Phe Glu Lys Asn Leu Glu Glu Cys Ile Thr Arg Ile Thr Asn Thr Glu Lys Cys Leu Lys Glu Leu Met Glu Leu Lys Thr Lys Ala Arg Glu Leu Arg Glu Glu Cys Arg Ser Leu Arg Ser Arg Cys Asp Gln Leu Glu Glu Arg Val Ser Ala Met Glu Asp Glu Met Asn Glu Met Lys Gln Glu Gly Lys Val

<210> 4990

〈211〉 589

<212> PRT

<213> Homo sapiens

<400> 614 Met Asn Asn Arg Lys Glu Asp Met Glu Ile Thr Ser His Tyr Arg His Leu Leu Arg Glu Leu Asn Glu Gln Arg Gln His Gly Val Leu Cys Asp ·Val Cys Val Val Val Glu Gly Lys Val Phe Lys Ala His Lys Asn Val Leu Leu Gly Ser Ser Arg Tyr Phe Lys Thr Leu Tyr Cys Gln Val Gln Lys Thr Ser Glu Gln Ala Thr Val Thr His Leu Asp lle Val Thr Ala Gln Gly Phe Lys Ala lle lle Asp Phe Met Tyr Ser Ala His Leu Ala Leu Thr Ser Arg Asn Val Ile Glu Val Met Ser Ala Ala Ser Phe Leu Gln Met Thr Asp lle Val Gln Ala Cys His Asp Phe Ile Lys Ala Ala Leu Asp Ile Ser Ile Lys Ser Asp Ala Ser Asp Glu Leu Ala Glu Phe

Glu	He	Gly	Ala	Ser	Ser	Ser	Ser	Ser	Thr	Glu	Ala	Leu	He	Ser	Ala
145					150					155					160
Val	Met	Ala	Gly	Arg 165	Ser	lle	Ser	Pro	Trp 170	Leu	Ala	Arg	Arg	Thr 175	Ser
Pro	Ala	Asp	Ser 180	Ser	Gly	Asp	Ser	Ala 185	Ile	Ala	Ser	Cys	His 190	Asp	Gly
Gly	Ser	Ser 195	Tyr	Gly	Lys	Glu	Asp 200	Gln	Glu	Pro	Lys	Ala 205	Лѕр	G1 y	Pro
Asp	Asp 210	Val	Ser	Ser	Gln	Pro 215	Leu	Trp	Pro	Gly	Asp 220	Val	Gly	Tyr	Gly
Pro 225	Leu	Arg	Ile	Lys	Glu 230	Glu	Gln	Val	Ser	Pro 235	Ser	Gln	Tyr	G1y	G1y 240
Ser	Glu	Leu	Pro	Ser 245	Ala	Lys	Asp	G1 y	Ala 250	Val	G1n	Asn	Ser	Phe 255	Ser
Glu	Gln	Ser	Ala 260	Gly	Asp	Ala	Trp	Gln 265	Pro	Thr	Gly	Arg	Arg 270	Lys	Asn
Arg	Lys	Asn 275	Lys	Glu	Thr	Val	Arg 280	His	Ile	Thr	Gln	Gln 285	Val	Glu	Asp
Asp	Ser 290	Arg	Ala	Ser	Ser	Pro 295	Val	Pro	Ser	Phe	Leu 300	Pro	Thr	Ser	Gly
Trp 305	Pro	Phe	Ser	Ser	Arg 310	Asp	Ser	Asn	Ala	Asp 315	Leu	Ser	Val	Thr	G1u 320
Ala	Ser	Ser	Ser	Asp 325	Ser	Arg	Gly	Glu	Arg 330	Ala	Glu	Leu	Tyr	Ala 335	Gln
Val	Glu	Glu	Gly 340	Leu	Leu	Gly	Gly	Glu 345	Ala	Ser	Tyr	Leu	Gly 350	Pro	Pro
Leu	Thr	Pro 355	Glu	Lys	Asp	Asp	Ala 360	Leu	His	Gln	Ala	Thr 365	Ala	Val	Ala
Asn	Leu 370	Arg	Ala	Ala	Leu	Met 375	Ser	Lys	Asn	Ser	Leu 380	Leu	Ser	Leu	Lys
Ala 385	Asp	Va]	Leu	Gly	Asp 390	Asp	Gly	Ser	Leu	Leu 395	Phe	Glu	Tyr	Leu	Pro 400
Arg	Gly	Ala	His	Ser 405	Leu	Ser	Leu	Asn	Glu 410	Phe	Thr	Val	lle	Arg 415	Lys
Lvs	Phe	lvs	Cvs	Pro	Tvr	Cvs	Ser	Phe	Ser	Ala	Met	Hic	Gln	Cvs	110

			420					425					430		
Leu	Lys	Arg	His	Met	Arg	Ser	His	Thr	Gly	Glu	Arg	Pro	Tyr	Pro	Cys
		435					440					445			
G1u	11e	Cys	Gly	Lys	Lys	Phe	Thr	Arg	Arg	Glu	His	Met	Lys	Arg	His
	450					455					460				
Thr	Leu	Val	His	Ser	Lys	Asp	Lys	Lys	Tyr	Val	Cys	Lys	Val	Cys	Ser
465					470					475					480
Arg	Val	Phe	Met	Ser	Ala	Ala	Ser	Val	Gly	lle	Arg	His	Gly	Ser	Arg
				485					490					495	
Arg	His	Gly	Val	Cys	Thr	Asp	Cys	Ala	Gly	Arg	Gly	Met	Ala	Gly	Pro
			500					505					510		
Leu	Asp	His	Gly	Gly	Gly	Gly	Gly	Glu	G1 y	Ser	Pro	Glu	Ala	Leu	Phe
		515				•	520					525			
Pro	Gly	Asp	Gly	Pro	Tyr	Leu	Glu	Asp	Pro	Glu	Asp	Pro	Arg	Gly	Glu
	530					535					540.				
Ala	Glu	Glu	Leu	G1y	Glu	Asp	Asp	Glu	Gly	Leu	Ala	Pro	Glu	Asp	Лlа
545					550					555					560
Leu	Leu	Ala	Asp	Asp	Lys	Asp	Glu	Glu	Asp	Ser	Pro	Arg	Pro	Arg	Ser
				565					570					575	
Pro	Pro	Gly	Gly	Pro	Asp	Lys	Asp	Phe	Ala	Trp	Leu	Ser			
			580					585							

<211> 30

<212> RNA

<213> Artificial Sequence

<220>

<400> 4991

agcaucgagu cggccuuguu ggccuacugg 30

```
<210> 4992
<211> 42
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: an artificially
synthesized oligo(dT) primer sequence
<400> 4992
gcggctgaag acggcctatg tggccttttt ttttttttt tt 42
<210> 4993
<211> 21
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: an artificially
synthesized primer sequence
<400> 4993
agcatcgagt cggccttgtt g 21
<210> 4994
<211> 21
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: an artificially
synthesized primer sequence
```

<400> 4994

gcggctgaag acggcctatg t 21

<210> 2001 <211> 2277 <212> DNA <213> Homo sapiens

<400> 2001

60 atactttagg ttataactta atgcaatgta ctttatattg ctgctcaaat tgtcccaggc geggeeceeg gaagetetet gggtagatee etgegeeett ggaegeeegg teetteagtt 120 180 ttltgagcac ctcaagcttc tggcctacaa aacgctcccg gctcagctgg agctttctgc geeegggtee tagagtegee cattleteta aggegeeitg geeetattit tagagagegg 240 tatttagaaa ccaagattag ggigctaaca atttittit aaattittat attittaaga 300 caggatetea etttgtaaca etteetttta gtggaagege egaceteetg ggagaeceae 360 geoccetgee geotteegte eegtttetea gaaaaceace cagacacece geoccaeegg 420 ccggggcccg ccgcgcatgc gcgccgaggc gtgacgtcag aacggcggcc aggacgccgg 480 540 acgtgcggca gttgcaggcg agcaggcgag gaatcgccgt ggcgtcttgg tgttctccac gctggttcgc aggtgaagag atggcgtttg tgaagagtgg ctggttgctg cgacagagta 600 660 ctattttgaa gegetggaag aagaactggt ttgatetgtg gteggatggt caeetgatet attatgatga ccagactegg cagaatateg aggataaggt ccacatgeca atggaetgea 720 780 teaacateeg caeggggeag gaatgleggg atacteagee eeeggatgga aagteaaaag 840 actgcatgct ccagattgtt tgtcgagatg ggaaaacaat tagtctttgt gcagaaagca 900 cagatgattg citggccigg aaattlacac iccaagattc taggacaaac acagcgtaig 960 tgggetetge agteatgace gatgagaeat eegtggitte eteaceteea eealaeaegg 1020 cctatgctgc accggcccct gaggtaggga gaaccctgag cctccagcag gcttatggct atgggccata cggtggtgcg tacccgccag gaactcaagt tgtctacgct gcgaatgggc 1080 aggegtatge egtgeeetae cagtacceat atgeaggact ttatggacag cageetgeta 1140 accaagical caltegagag egetalegag acaaeggeag egacelggea eigggeatge 1200 1260 tggcaggage agccaeggge atggeetlag ggtetetall ttgggtette taggggeete aaggletiga igigealage lielgalaac eelgigigea alaalaigal ligeagggea 1320 tttctgtttg tgacaaaagt ttttaataat agttttaatc attcctttga aagtagtgat 1380 gleataattg tactaateea cataagtace acagagaagg gittgaactg tgctatittg 1440

ttcaaatgtt	gactctccgg	gggcactggc	tcattccaag	actgttcttg	tgcaactctc	1500
agaatacctt	atttgagcat	acctgttttg	aaaggcattt	tctttttaga	gttaggtgta	1560
gtgcttaagg	gttaatttat	tttcatgtta	tgccagtaat	atagtgttgt	atgcctattg	1620
agtgattgtg	gcaagaaaaag	ctacagette	tttgcgttta	actititcaa	accacagacc	1680
agaactggtt	gcatgttact	ttaggagttg	tgggttggta	agctcccagg	tacttcccga	1740
ggctatggtg	tgagagcccc	cgtcctgccc	tctggggctc	cacaggcccc	tggcaaggcc	1800
gatggctcag	gatgatgggg	cacageeege	ctttgaacaa	tcatgcttca	gaaatctgcc	1860
tgaccctagc	tgctgctgct	gctcacttta	ttcttgtatg	gctttggtag	gcatacttgg	1920
agaacatatc	ccacattagg	aattgattta	agcctgagag	tttgagggct	ttaatccttt	1980
aaaacttgga	gaagctggct	gggcgcggtg	gctcacgcct	gtaatcccag	cactttgaga	2040
gaccgaggcg	ggcggatcac	gaggtcagga	gatcgagacc	atcctggcta	acacggtgaa	2100
accccatctc	tactaaaaat	acaaaaaatt	agctgggcgt	ggtggcaggc	gcctgtggtc	2160
ccagctactc	gggaggctga	ggcaggagaa	tagtgtgaac	ccaggaggcg	gagcttgcag	2220
tgagccaaga	tagtgccact	gcacttcage	ctgggtgaca	gagtgagact	ctgtctc	2277

<211> 2276

<212> DNA

<213> Homo sapiens

<400> 2002

60 ctatagattt tatgaateec ategttacat ateceacite agtaggieti gggiggeeca agactatgtg ttaacaagtg gitettaige aagitgagaa acaetggett alatagaeea 120 180 aatottgaaa actgggtata tacattgtoo gtaatgagag agtgccactt collgccaat 240 accetggtat tatatggeeg attitigtete titigeeaata attiteatiat aaacigitea 300 getgtgttga ageaaaactg tagaaaaagt eetgtettea teagatttte tgaggttgta 360 attatactet igicatacca giggagacce agiaateata eigeaacaat igigiaacac 420 ttgcattica tactcaggca aaacccagtt ataaaggtag citcliccic alltitggtt tttccttcac ttttagaaag tacttagcca gtagttcttg cattatttgt ataaggggga 480 totgtgatgg cagcaggatt attactgata tataaagtaa gttttattot aagatotatg 540 ttacaaatti totattgigg gaaagagaig ttagaaccag aaciilgggg alagcaccaa 600 660 agatactaga aaacagacat ttataaggta tettttttee eeetettta ggacatgaaa 720 tetgetgtga teacgeettg eagleatitt ticcatgeag getgtettaa gaaatggetg 780 tatgtccagg agacctgccc tctgtgccac tgccatctga aaaactcctc ccagcttcca 840 ggattaggaa etgageeagt tetaeageet eatgetggag etgageaaaa egteatgttt

caggaaggta	ctgaaccccc	aggccaggag	catactccag	ggaccaggat	acaggaaggt	900
tccagggaca	ataatgagta	cattgccaga	cgaccagata	accaggaagg	ggcttttgac	960
cccaaagaat	atcctcacag	tgcgaaagat	gaagcacatc	ctgttgaatc	agcctagagg	1020
agaagcagca	ggaatgatgc	tttgatactc	tggaggagaa	gttaactcaa	gatggaattc	1080
atgttctgat	ttgaggaatg	aaaatgagat	gatcaggcag	gaaactgaca	ttccaaggat	1140
ctaatccagg	aagtactctc	agtggggacc	acctgctttc	atcccctgac	attgtgggag	1200
aaattttgca	atgtatgcta	atcaaaatgt	atttatatgt	tctctgctga	tgttttatag	1260
aggtttgtga	agaaaattca	acctcagcaa	cttcagaaac	tgcccctgat	acgtgtgaga	1320
gagaaataaa	atcagatttt	gagtgttgaa	gggactgagg	aagtgaggat	aaagagcatg	1380
aggacagcat	ggaaagaagg	aggcagaagt	ggaactgaac	tttcactctc	catgggacag	1440
atcaatctca	ttatcaagtc	tgaatagcaa	ccagccctct	cctccacccc	gtttctcctc	1500
agttaattgg	agctcagtca	ggtgattatt	gagtcttgta	cagcactgaa	atgaaatcaa	1560
agatgaagaa	gcattgattg	tattcaaaga	ttgaagcacg	ctcatacttt	gtatgtgctt	1620
tagggaaggg	gtgggtgggc	acttgggcct	tgcgggtgca	ttcatgtaat	ctgagactct	1680
tgaactttat	gacggagtct	tcagtatttt	gatgtatatg	aaacttttgt	taaatatgtt	1740
gtatacttcg	ctggctgtgt	gaagtaaact	aaaactctga	tgaacacttt	ggagtctgct	1800
ttagtgaagg	agaccaaagt	gggaagggct	ttagggcact	gatagaggcc	ctgggtgtac	1860
ttttcaatcc	tgtgtaatgt	ttaattcttg	caactgaatc	aaaacagtgt	taaattatgg	1920
caatatttgc	actttgggaa	tgaatacata	actgtatgat	cacactctgc	aaatgccact	1980
tttaaagctg	ttaatagact	ttgcaccttt	tctttgacaa	ggatgtgtca	tatttaaatt	2040
tttacactca	tcatggctac	aggtagaact	ggggaggggg	gaatgtaatt	ttttatggga	2100
attttgatat	gaaaagaaac	tagtcattta	tttatacaat	aggcttggct	caaaaagtgt	2160
ttttcagacc	tcggtattcc	taatgtggga	tgtgacttta	ttttatttt	agtagcaaat	2220
ttggatgtag	actgacagac	acagctgaat	gtcttaataa	atttaaattt	gaagat	2276

<211> 2076

<212> DNA

<213> Homo sapiens

⟨400⟩ 2003

cacactgagg ggacagtetg gaggettgca gtgacteaga cacagceaat teeteecta 60
atageactga ateaeggtte cageggecag tggtegeece tegteaaggt etaaggetge 120
tgeageeceg geteeeggag geegttteeg egegeacaeg egeateeata egtacagaeg 180
tgetegggat gegggteeeg eeggeggta eetgggeaet gegeeceate tggactgaaa 240

```
tggggacacc ccttcggggg tcccaggctc ctggccgtat tgttctcctt ctcctcgtga
                                                                     300
taacteegea gtggaggtgg atteegteea agaegeecaa egtggeteeg egtageaate
                                                                     360
agegetgeaa teetggeggt taceteageg geggegtete tetetgegee teacactege
                                                                     420
agcccgcggc cclccccaac ttagggcgtt tacaaaagaa actactccag acgcgctgca
                                                                     480
aagggaggcg catgtgccg aaagctggcg atcagacggg gggggcattc tgcatgtgtg
                                                                     540
atgittetgg gggcggtggg gagtgtgtgt eggggteggg gggcgggggg gagteaggea
                                                                     600
gaaagacagg gacaacctcc gctatgaagg atccgcgagt cctcaaatgt aagctccgtg
                                                                     660
tgactaacga cctgcactga tttggagagc gggcatgtta aaggtcacgg acaattgttg
                                                                     720
ctggcttcag catgaatgcc taagtgggat gtattcttca gcaatcacgt ttaagtctga
                                                                     780
ttcaccgaaa agtattgacg tgcccaccat tcatttcagt acactgtgaa aatgcacaaa
                                                                     840
gaaagtatcc ccaaattcag ttaattacaa agccgtaaat gtccttgtat acacatatta
                                                                     900
ttacatacat gtaggtaaca acaaagatta aaatttgaag acactttaat agctttttgg
                                                                     960
taggattitg gaatgaatat cagteetgta aacctaegit catetgeatt ettgggteta
                                                                    1020
ttttaaagta caaacttgcg ctaacaattt ccatgtgttg aaaatggaca aggtagatca
                                                                    1080
ttgaatggtg atcaagactt ccaaacccct ccacataaaa ctgttcatga cttgcttcct
                                                                    1140
ttttctagcc ggtttagggc cctgtcttaa gtcacccaca tgtgatttca ctcagggcat
                                                                    1200
tgtctgtcta caataatatt gtgcttttaa accatttcct ttcttacacg tttatctaca
                                                                    1260
gtgcatgcga aatctgagag cgtaatttga tggatgggca aagagttaag tcctggtgtc
                                                                    1320
tggtgtggca gacctagaaa atggcagctg gagggccagc atcattttgt tactgacaat
                                                                    1380
tgaaacgtgt tcacattgat tgtacacaag tcactggtgg ttgttcattt gtcaatgcac
                                                                    1440
tattcctagc tcactccaca cacacaaaaa aggtataaaa atcaaatgtt taatacaagt
                                                                    1500
ttecataeta tteetglaac catatttage attgecaaca ttteaactgt tttaataget
                                                                    1560
teaaacactt aaagtaacca ttagggatta agggcaccgt ttgcccctgg aatggcccag
                                                                    1620
gagagettet ectattitga aaggittaeg taaattatag tattiggatg gageaaagte
                                                                    1680
agcagtatta atggttgaat attaatggtt gatttiggct actigiitta itilagigat
                                                                    1740
atgtgatatt ttacacatgt atggggtacg tgtattigtt acaagcgtag aatgtgtaat
                                                                    1800
gatcaagtcg gggcactlag ggtactcatc agctgggata titaligiti ctalgcgitg
                                                                    1860
ggaacattic aagiicigic ticlatetat titgaaatac acaatccatt gitattaact
                                                                    1920
gtagtcactg tagtctgcta tcaaatatta gaactactcc ttctatctaa ctgtatgttt
                                                                    1980
gtacceatte actaacttea ticcecceca cectetatit ataatiitat aacagacaat
                                                                    2040
aattttggtt aatgaaataa atgggggaaa gaaagc
                                                                    2076
```

<211> 2525

<212> DNA

<213> Homo sapiens

<400> 2004

ggcctttttt ttttttttt tttttttgag atggagtctc actctgtagc ccaggctgga 60 gtgcagcggc aagateteag eteactgeaa ceteegeete eegggtteaa gtgattetee 120 tgcctcagcc tcctgagtag ttgggatcaa tcacaggcac gtgccaccac gccctgctaa 180 tttttgtatt lliggtagaa atggggttic accatttigg ccaggctggt ctccaacttg 240 tcacctcagg tgatctgtct gcctcagcct cccaaagtgc tgggattaca ggcgtgagcc 300 actgcaccca gccatggtgg gtgttttgta gggaacaatt tcaaaaggac ttctggtggc 360 420 aaccattgag cctctggttg acagatatgg gtaaaattat tcagaaaaca tatctaagac aggatgtgga gaatagtact gtcatcagtt tataccttaa taccacatct aacaatgttt 480 atgatagggt tgatcacttc catgaaggca tcacaagcct tgctgtgtga agggcatctg 540 600 aatacatttt aatattilat alcigitett cacacettag eeecteacte iggagaaaat aglacatttt cittcilaaa ataiggiaca citaagccic aaaigiggai citticitig 660 aaagtaaaac tgaacaggtc cttctgccca cctgcagtcc ccaaggaaag aacacatgtt 720 acgitcatig ccaalaatag giccitcagi actigitgaa igaagaatac tigigiiitt 780 ccactggcca accaaggtgg atcctgaaag tggaacccgg agttctctaa taactaaatt 840 agtgttttta gtagcicatt tigaatccci aagcigtgac ticaacicig aaaggciggc 900 taactctggg aggttacctt cacttaatta agtacagcat ttcttccaaa gcgcatgcag 960 tgctttatgt aaattctctc tcctggattt gtgtgacgta gcagggttag aatggtgaga 1020 1080 cagatgcctg gttttggagt cataagactg gctttgccac ctagcatctg tgtggtctta ggccagccaa cttttctttt tttttgagat ggagtctcac tctgttgcca gattggagtg 1140 1200 cagtggcata atctcggctc actgtgcaac ctccgcctac tgggttcaag tgattctcct geeteageet eccaagtage tgggattaca agegtgagee acegeaeeeg gecaaagata 1260 cgtttttaat aactiggget ettteaagag aaacagggag caccateace teagaaagee 1320 tttaccacte actgctgccc caaaacaaga gatgcatata tigttgacaa ccagtgcttg 1380 aattaattac attttaaaat atcgtcctga gctctgcctg tagctgagag gctgagaagc 1440 gtgaaatagc caggattaaa tgacctgcaa atctagactg gcttcttttg gggctggtac 1500 tgccaggcag acagateeet gtteettgca ecceeactgt cetecaccat etetactetg 1560 gatcaagggt caaaaaactt ttttttgaga tggagtcttg caggctggag tgcagtggca 1620 tgatetegge teactgeage eteegeetee eggetteaag eagtteecet geeteageet 1680 cccgagtage tgcgactaca ggtgcacacc accaegeceg gctaattttt tgtagtttgg 1740 tagagacagg gittcaccat gitgitcagg atggicicga iciccigacc icgigatccg 1800 cccgccttgg cctcccaaag tgctgggatt tagaggcgtg agccaccgcc tctggccaca 1860 1920 ataggaaata etttaggett eagggeeate eagtetetat gteaactaet eaattetgee 1980

ttcgaatctg	aaagcagcca	cagataatac	aaacacaaat	tggtctgggc	tgtgttccaa	2040
taaaacttta	tttacaaaaa	caaatggcca	gccccaaggg	cctggtttgc	aactcttgct	2100
ctggagcaga	gcagaaggta	tactctgaac	tgcaacaaag	tttctgctgc	aaaagcagca	2160
cctctgctgt	ccgtcccctc	ctctctgtcc	actggctctg	gacgtccatg	tgaacaggct	2220
tgccaagaag	gacaaagtgg	gcaggtaaag	ctgggggggg	cggccacaat	caagatccca	2280
acacccctat	ctttaagagg	cagtgccaag	cgaatcccat	ttcaggggac	ccactctacc	2340
tcgctgccta	cgatgaattc	ccatcttaca	gcctctcgat	tactatgcag	ttaccaagct	2400
ggctaccacc	ttactaagat	tcttgccatt	ttctcattct	agtcaaaaaa	gtaagtcatc	2460
ggtttagtgg	agggggcagc	taaagcccaa	gtttgtattt.	gagaaagatg	tacaacaggt	2520
tcttt						2525

<211> 3574

<212> DNA

<213> Homo sapiens

60	tccctcacca	ccaaaagggt	ccatgaacac	gagagggcag	tctggctacc	acatctgttt
120	gtcccccggg	gggttccctc	gcatccaaag	ctcgccatga	cagaggttcc	taaacgtcca
180	gtgaagttcc	tccggctgag	tgcgggtgat	gacatgcagc	ctcgggtaga	atatggatag
240	atttgccgct	tgtgcataat	tcccgattac	gtgtaccgcc	ggccgggagg	tggacacgat
. 300	atgttgacca	gttcaaactg	tcaagccaca	aaggagcccg	aatccgattt	ggaaccagaa
360	tatcatcctg	tatggtggaa	agatgacagc	tctggccttc	agaacttgct	gtctggataa
420	acaacagaaa	agaaaataaa	ttatttcaat	gaccggctac	agacactttt	ataaagacga
480	gttaattttg	tgaatcagta	aattggaaat	ccatcctgtc	tgggttgatt	ttcctctaat
540	aaccatggca	tactatcact	ctaaagagat	aaagtatatt	tgccaatagt	gcacactggt
600	atttttccaa	acccatcctc	acggccaatt	gcagaatacc	catatttaag	aagctccagg
660	tgtgcagacc	agtagatttc	tggttattaa	aagtcatcaa	cgtggatgct	ctagtggtat
720	gagatgctct	aggtcaacct	tgattttgca	gaggcaatag	tgtagatgaa	agccaagaat
780	agtagtgaca	attaagcatg	ttattgaatt	gttgagcaga	agctcatatg	tgagtatcaa
840	attaaacatg	atcatcaaaa	ttttcttcgg	tttggtcctg	atgcatacac	gaaggctgga
900	caagatgatg	ggccatcata	taaattgggt	ccagagccca	caataatagc	cacgtgtata
960	ttaaataatc	agatattgct	aacaaagaac	acagatattc	agaattgggt	ccgtgggaga
1020	tgtcttccta	cattatctcc	atactactat	aagaacatag	aagaaaaata	tcacctacat

```
atgaagggac titacaacci tatcaaaaga cigtaattac attitgitte accccaaagc
                                                                    1080
taatggctgt tggtaaaaag gatattggac cttcatacag acaggactat gctctcttt
                                                                    1140
tgagattiga giccgtagga agiaaagatg gattittgag agatgatgac tataaaacca
                                                                    1200
tcaaaagtga acgatttcag aaagtggaat tagcactgac aggcacagga cttcctgttt
                                                                    1260
tactacagtt igatceagga ceagitetta attitaaace tigitteatg ggigaacgit
                                                                    1320
cagaaattca gtgcatcata aaaaatcaat gcgaattact tcctgtgacg taccacttta
                                                                    1380
aaaaaaactgc aaattitgaa attgaleetg aaaagggcaa gattactgga gggggtatgg
                                                                    1440
tggatgtgat gtgttcattt gttccacatc aacttggagt cttcaaagtg aagcagatga
                                                                    1500
tagagattat tggtttagtg gcagaagaag atttgcaatc tttgtcggta aaatctttcc
                                                                    1560
atcacgtata titagettic aacagcatet giaaaactic caccaagaaa giigtgatga
                                                                    1620
aatttgatcc tggtatattg ccttcgatcc gtaatcccac gggaaagttt gtggtcaaag
                                                                    1680
acttggcaaa acgcaagaat tatgcacctg tagcaatgct tcaatcagcc atgacacgca
                                                                    1740
ctcacaatca tcgctcatgt gaagagccag tgaaggatat gctattagcc tttcccaatg
                                                                    1800
accgagetge aactateagg tetaaagace ateataaaca titeaggeea attiteacaa
                                                                    1860
augticcaug attiaaciai gigaalcaig attitigcata tactacatti gaaaaacagc
                                                                    1920
aaaagaaatt acatgaaaac tattatgcaa tgtatctcaa atatttaaga agtgtgcgct
                                                                    1980
tgcagaagaa acaagcagag agggagcgca tgtattcata tgatgataca gacataggct
                                                                    2040
tagagccagg atcaggteta aagtcaccet cactetcaga ageggaaata gaagaggage
                                                                    2100
tgtcttcagc agcaaattca attagagcga atcgattgtt aaccaccagg ggtatagcat
                                                                    2160
ctcaggagga agagtctgtg agaagaaagg ttctcaaagg acttaaatca gaaccatcca
                                                                    2220
ciccacaaga aaaacatgat tgcagciiaa tgligacacc aaagcaaati catcaagtaa
                                                                    2280
ttgttgggcc ttctgtcctt aacittggta atatttgtgt gaacictcca aatactcatc
                                                                    2340
tacticatgt tattaatatg ctacctatgc atgittigct ccagttagat actgattag
                                                                    2400
aagaacttca gaagaccaac caattttcat acgtgattct acctacatcc agtacttata
                                                                    2460
tttcaatggt attigattct cccaccattg gaaaattttg gaagtctttc acctttacag
                                                                    2520
tgaacaatgt acccagtgga cacatectag tggtggcagt tgtccageca gtaacacttg
                                                                    2580
agctatette taatgageta gtattgagae caegaggett etteatgaaa acatgtttte
                                                                    2640
gggggacagt tagatigtat aaicgicaga aiigtigtgc icagiitcaa iggcaacccg
                                                                    2700
taaacacagg aagagggata gcattitcia titgtccatc taaaggcact gitgaagcat
                                                                    2760
attcctcact ggaatgtgaa gtaacttggc agcagggctt cagttctcca gaagaaggag
                                                                    2820
aatttattet teatgtettt caaggaaaeg egitgaaget aaaatgigit geacatgtaa
                                                                    2880
ttattttcct tgaacatggt tittgttttg agggctatga attggttggg tatacactgg
                                                                    2940
tgtatatagt tacctatatc tagaattaac tgtaaaaccc aagactttca tgcaacagta
                                                                    3000
ctagtitttt tgttagagcc tctataaata tgtaatatca tcatgggagc cattgaaatg
                                                                    3060
aaattatttt attaagagac acaaaaagta ttttcagaga atalacttga tggattaaaa
                                                                    3120
algigagiag agggaaagci glaatalgca allitaacci tiltciggia cagiccagag
                                                                    3180
```

g	gccttaaat	tcatgactca	atcaccaagc	atgattttac	atgtgtacca	aatttcccac	3240
t	caatgttct	tagaaatatt	aaagaagcca	aatgctcttt	tactaaaccc	catctatatt	3300
t	ctaggacat	gatgatactc	ttacatattt	cagctgtgga	ggagtttta	gcctcaagag	3360
a	tgagaaatt	catctacttt	tagtgatggc	aagtgacaga	actcagtatg	gtttttcttc	3420
t	aagcctaaa	ataagctggg	tcctactact	tttcattatg	tgtaaattag	ttttattttt	3480
t	aaaaacttt	ctattgaagt	ataacatgca	tatgtatatg	tatatgtgga	gaaacatgaa	3540
g	tgattaaat	aaaatattca	tttgtttgtc	attc			3574

<211> 4634

<212> DNA

<213> Homo sapiens

<400> 2006

attgagctgg gctgcagagg agtgtgaggt gcagacacca tgaggtaccc acagccagga 60 aaacgaggat ggtcggggag acgcgccagc gaagagctga gcccctgcgt gggacccctc 120 agtggttccc agggggcgtg ggacttgcgc agtcctttca gagggctgtt taccaacagg 180 aaccgtaaca ttaaacctgc tcagacccct tgactcagca atttcatgtc tgggaatata 240 300 tcttaggaaa ataatcagag atgcctacca acatatgtga tgatgatgta tgacagaatt 360 attatacaaa tatatccata giaacagggg gitigcigaa ataaattaic atatattcat ataatatgac altatcaggc cattaaaaat cacagtttca aagagtaata aaatgggaac 420 480 atgctcatag talagittii taaaaitgca gatggtatai ggctaaaaat gictaataat gcaaagatgi alacagacci taaleeleta gceteeteee tagagatgae etetgitaat 540 tictcaaata tittictgga tactitacac actcacacac tittittigag acagagtitc 600 actittgtea eccaggetgg agtgeaatgg tgtgatettg geteaetgea accteeaect 660 cccgggttca agagattete etgecteage etcccgagta getgggatta caggtgeetg 720 ccaccitgce iggetaatit titgiatitt tagtagagae ggggitteae cacattggte 780 aggotggict caaactooig accicaggig aloogooigo otiggocioo caaagigoig 840 ggattacagg cgtgagccac tgcgcccggc cattcatcit aaittitaaa aaatctaacc 900 atgaageett ggttatetig gagagettie etgattagea caaaaagaaa aaaaaateea 960 attetttaca getgeatact attecattat tigtatgigt catatttiat tiaaccatee 1020 tgctattagt gaccattgag tiggcttcct gtgttttgcc gttacatggt tgcaacaaac 1080 atgittgcat gigtcigccc tcatgigcat gatacaigat igatitigata gatittagga 1140 1200 atlacateat teatleatae acteageaaa tatttaatga gtgeetaete tetgataggt gctgttggat gtggctaaat tttaaagtgt agaatttaaa aggtggctac caaattccat 1260

```
gtgcaaaatg accccacgca tgtataaaaa cacacactc cacagattta tatgcgggag
                                                                    1320
agaagatgtg gtccctggcc tctaggctct ctcagtctgt ggcaagacag acagacatgt
                                                                    1380
gcacgcggca cigiaaggii gagcacagic taagtactca gcatggictc tggcacatag
                                                                    1440
taggtgccca agaaatacat gtcgaatgaa ttgagggggt aaggccttct agggcaggtg
                                                                    1500
geeletgace teageettea gtgtteegta ggtggaatta tetgeeagag aegtggeaaa
                                                                    1560
agggagagga accaagactg aggcacagag gitcaaacgi acccggcaca itcagagaat
                                                                    1620
ccttttcaga atcacgtccc caagagcttc tgtgttctgt acggtgatgt tgcagtgctg
                                                                    1680
tiliticegea gictegetee aleggeetea aleegeigta calcatgetg eeeigtaeee
                                                                    1740
tgagtgcctc ctttgccttc atgttgcctg tggccacccc tccaaatgcc atcgtgttca
                                                                    1800
cctatgggca cctcaaggtt gctgacatgg taacacagct gtttttattt actcccgtcg
                                                                    1860
gactataacg ctgttgtcat aagggatgcc ccatttatga atgacagagt ttcaaaacga
                                                                    1920
tglcatgtga cttgggaatg ccacggaaca tccagacctg tagccattgt tgacatttat
                                                                    1980
aatgeagett ttettetttt tetgagatga teteaageet eacacaetgt tettteteig
                                                                    2040
aggigggila tagacicice cacciggaga agccigigea ggcaccaggg gagicciigg
                                                                    2100
aaggggtgaa ggtggggctg agggactcat atggccaagg atgaacttga caaattagca
                                                                    2160
agaaccatga agataggcag ggcaggctta ggcagcaggg ggatgttaat gacagtcaca
                                                                    2220
gagattigta ggggtgcctg aagaggtaga agcagggaga gggagagaga gagcactgcc
                                                                    2280
tgggagtaga tgatgccttg gaaacaaatg tagtcagagg aagaactctt cattagctct
                                                                    2340
gtcacctttg ctgggagaag ggcagctttg cagctctggg ctgggaaaga ggcaagtgtt
                                                                    2400
tgagcccaag aggccagaaa tgtacctggg accaatcggg tgttcgttat ctcagagcct
                                                                    2460
ctgctgggta. tctcagggac tccatgagca ttttcaaaaa aaaggtgggt cccagaaacc
                                                                    2520
atggactgca aacttgactc caatccccag taaaatatct acaacagggt agtgaagcga
                                                                    2580
tggttagtga ccatgaggga agcttgcaga gcaggcatca gaaagagcct gaggaggtcc
                                                                    2640
acagggaage tggcacgice itgiaggata gitaaggcac tggggtgage aatgaaceig
                                                                    2700
gacteaegga acactggget etgtgaeegt tteeetgaat ggeetaaget gttgeeteet
                                                                    2760
gteacticic tgaggicatt ticcaaatge geaegggeat agagaaceea teeacteige
                                                                    2820
ctacttccca gggatgcctt gagcactgag gatacctggg ggacatgaag tcgcactgtc
                                                                    2880
ctgggggtcg ggacacccca gccagggaca gagcatggca cagggacatc gaggcccagt
                                                                    2940
gagccgaccc tttgtcctcc tctctgagag cactagtccc cagcaggcct cagggtgctg
                                                                    3000
actetgtete titteeaggi gaaaacagga gicataatga acataatigg agicticigi
                                                                    3060
gigitiitgg cigicaacac ciggggacgg gccatatiig actiggatca iticccigac
                                                                    3120
tgggctaatg tgacacatat tgagacttag gaagagccac aagaccacac acacagccct
                                                                    3180
taccetecte aggactaceg aacettetgg cacacettgt acagagtttt ggggtteaca
                                                                    3240
ccccaaaatg acccaacgat giccacacac caccaaaacc cagccaatgg gccaccicit
                                                                    3300
cciccaagee cagaigeaga gaiggicaig ggeageigga gggiaggeie agaaaigaag
                                                                    3360
ggaacccctc agtgggctgc tggacccatc tttcccaagc cttgccatta tctctgtgag
                                                                    3420
```

```
ggaggccagg tagccgaggg atcaggatgc aggctgctgt acccgctctg cctcaagcat
                                                                    3480
ccccacaca gggctctggt tttcactcgc ttcgtcctag atagtttaaa tgggaatcgg
                                                                    3540
atcccctggt tgagagctaa gacaaccacc taccagtgcc catgtccctt ccagctcacc
                                                                    3600
ttgagcagcc tcagatcatc tctgtcactc tggaagggac accccagcca gggacggaat
                                                                    3660
gcctggtctt gagcaacctc ccactgctgg agtgcgagtg ggaatcagag cctcctgaag
                                                                    3720
cctctgggaa ctcctcctgt ggccaccacc aaaggatgag gaatctgagt tgccaacttc
                                                                    3780
aggacgacac ctggcttgcc acceacagtg caccacaggc caacctacgc ccttcatcac
                                                                    3840
tiggiteigt ittaategae iggeeectig teecaectet eeagigagee teetteaact
                                                                    3900
cettggtccc etgttgtctg ggtcaacatt tgccgagacg cettggctgg caccetetgg
                                                                    3960
ggtccccctt ttctcccagg caggtcatct tttctgggag atgcttcccc tgccatcccc
                                                                    4020
aaatagctag gatcacactc caagtatggg cagtgatggc gctctggggg ccacagtggg
                                                                    4080
ctatctaggc cctccctcac ctgaggccca gagtggacac agctgttaat ttccactggc
                                                                    4140
tatgccactt cagagiettt catgccagcg titgagetee tetgggtaaa atetteeett
                                                                   4200
tgttgactgg ccitcacagc catggctggt gacaacagag gatcgttgag attgagcagc
                                                                   4260
getiggigat eleteageaa acaaceeetg eeegigggee aatelacitg aagitaeteg
                                                                   4320
gacaaagacc ccaaagtggg gcaacaactc cagagaggct gtgggaatct tcagaacccc
                                                                   4380
cctgtaagag acagacatga gagacaagca tcttctttcc cccgcaagtc cattttattt
                                                                   4440
ccttcttgtg ctgctctgga agagaggcag tagcaaagag atgagctcct ggatggcatt
                                                                   4500
ttccagggca ggagaaagta tgagagcctc aggaaacccc atcaaggacc gagtatgtgt
                                                                   4560
ctggttcctt gggtgggacg attcctgacc acactgtcca gctcttgctc tcattaaatg
                                                                   4620
ctctgtctcc cgcg
                                                                   4634
```

<211> 3576

<212> DNA

<213> Homo sapiens

ggggaagggg	aggaggaagc	caccctgtag	acttgagact	gagtcttaat	tcaagttcaa	60
actctgttgt	taaccaacat	ccaaagttat	gcaatagctt	acactgcctc	tgttaaaaaac	120
tigigaaata	tcactcattg	ataaactatt	gtaatacttt	tccttagctc	ggtttctcaa	180
ctgaggcact	gttgacattt	caggccaggt	aaccctctgt	tttaggggct	gtcctgcgca	240
ttacaggatt	ttagcagcat	gcctggcctc	tgcccactca	gtgccagtaa	caccttcctc	300
agcaattcat	tacgtctgtc	agaaatgtct	ccagacattg	ccagatgtcc	cctggagggg	360
cacagttgcc	tccatttgag	aggccctgct	tcagaggatt	cactctgagt	gagttcgcta	420

atgcatttga	gcaaattgga	agttcttccc	tgggccagag	gctcagtagc	caaaacagaa	480
ttacccagag	aactaggcct	ccgtagaaca	gtcattgcct	gaaaggggca	ggaggtgact	540
gggcggaatg	gcacaagtgg	ccccagagca	ggtccagccc	cctcccaccg	cagcatccag	600
aaagacccgt	gggcattcgg	tagatgagcc	caagatctag	aaatggaaca	ttactggaga	660
aaagggccta	ggagactaga	ggtagctcta	ctctcagtgt	gagcgtgtgt	cagcacaggc	720
gttgtggtgt	ctgatcacag	agtaaaggta	tgcttcctta	atcttgcatt	gaaaaccatc	780
tecttegeat	acaccatatg	caaaaccaaa	ttcaggtaga	ttaaaaaagcg	agaaaagtaa	840
acaaaactgc	agatgcattc	aggataaaag	taagataata	attttattgt	gttgagttat	900
gaaaagcctt	ccttaaaaaag	atacagccca	gagatgagaa	aggaaaaaggc	acaaaaggcc	960
cctgtcatgc	gccatggatg	aagatacaag	ttgaatgcca	gaaagcgagg	ggcacaattt	1020
aaagtgttca	tttttagatt	tagcaagtct	actttcacac	atgtatccta	taaaaatatt	1080
tgcacatatg	cataacggca	catacaagga	cataactgca	gcaatggcaa	ggagtgatga	1140
aaaagtagga	acagtggcca	aatcgagtga	taacagaaaa	ggaggcagca	ctgtgaggaa	1200
ggttgcgcag	agtgcaccgc	tgagcacggc	ctgcgcctag	acccctgtgc	tgtctgagac	1260
cacctctgga	gtatgcagcc	atgtgtggat	cacaggtgtc	aaatagcgaa	gttactctgg	1320
aagagtttt	tttgtttgtt	tttttggggg	gttttttgt	ttttttttg	ttttgttttg	1380
tttgtgcaga	cagagtctcg	ctctgtcgcc	cacactggag	tgcagtcacg	tgatgtcggc	1440
tcactgcaag	ctcttgcctc	ccgggttcac	gccattcgcc	tgcctcagcc	tcccgagtag	1500
ctgggactac	aggcgcccgc	caccatgcct	gtagtcctaa	ttttttctgt	tttttagtag	1560
agatggggtt	tcaccgtgtt	agccaggatg	gtcccgatcg	cctgacctcg	tgatccgcct	1620
gcctcggcct	cccaaaatgc	tggaattaca	ggcatgagcc	atcgctcccg	acttaatttt	1680
gcattcttag	tggagacggg	ggtttcacca	tgttggccag	gctggtctcg	aactcctgac	1740
ctcaggtgat	ccactcgcct	cagcctccca	aagagctggg	attacaggtg	tgagtcactg	1800
cgctcagctt	aattitgtat	ttttagtaga	gatggggttt	ctccgttttg	gtcaggctgg	1860
tcttgaactc	ctgacctcag	gtgatccacc	tgcctcggcc	tcccaaagtg	ctgggattac	1920
aggcatgagc	cattgtgccc	ggccacattt	ttctttttaa	atcattttta	ttcaggtaca	1980
acttatccaa	aaatcagcac	cactggtttg	tttattgcag	aaaaatgaaa	tttagaagtt	2040
tggtctaaat	tttctagctc	gctaaggaat	cttcgaaaat	tcccaatttt	cctatttctc	2100
actaatgtag	gaaatattta	aaagccagca	aagaagaaaa	catctttaa	aatctcatta	2160
tctatacgta	atcactaaga	accttttgca	actttccctt	atagttttt	aacctgtata	2220
tgaggcgttc	tctgtcctga	agtaatgtcc	tgcctctggc	tagctcctgt	gacggtagcc	2280
ctcccggggc	tggccctggg	tgaggagggg	tggcggcggg	gaggtgagcc	caggaaaggc	2340
tgccctcgcc	aaggctcgga	aacttcattc	gtgcaccgca	cgaggcgatg	gctcagggca	2400
ggcttggaca	ccaatacttt	gccagctcct	gaggcaccgg	acaggetetg	gccagagctt	2460
aattggttag	ccctagaacg	ttccacgttc	acgtcagact	ccatagtagg	gactttctcc	2520
tcagagctgg	gcaggaggag	cccactgagg	gtgtgccatc	tctgccctcc	agggaaagcg	2580

```
ggaagcaaca gggaaacatc catctgctcc gccctagagc ccctgtcaat tttggaccca
                                                                    2640
ccgctatagg tcttctgccc catactgtta gaaaaagatg caggttacct gggcacgtaa
                                                                    2700
acggttitca ggagiggagi gcctgagatc ccagagicca cctticctti atataacact
                                                                    2760
cgtgtcacag gacagattag atttcttccg tgtttggaga acattagtcc tttaaaatat
                                                                    2820
cagectgtgc tgcaaagtgg ggtggattet ctagtetcag teactgtetc ageagtgetg
                                                                    2880
ttgaageeet eteacetget eettetggae tteetaggge tgeagaceae aagaetggga
                                                                    2940
aaccactigg aagaccgagt gaacaaattt tigcggcgcc agaatcaccc igaagccggg
                                                                    3000
                                                                    3060
gaggitting teegagiggi ggeeagetea gaeaagaegg tggaggieaa geeegggaig
aagtcacggt ttgtggattc tgggggaaatg tctgaatctt tcccatatcg aaccaaagct
                                                                    3120
ctgittgctt tigaggaaat tgacggcgtg gatgtctgct tittiggaat gcacgtccaa
                                                                    3180
                                                                    3240
gaatacgget etgattgeee cectecaaac acgaggtatg tgacagggea catetgggee
                                                                    3300
tgicciccaa gigaaggaga igatiacaic ticcatigcc acccacciga icaaaaaata
cccaagccaa aacgactgca ggagtggtac aaaaagatgc tggacaaggc gtttgcagag
                                                                    3360
eggateatee atgactacaa ggatatttte aaacaageaa etgaagacag geteaceagt
                                                                    3420
                                                                    3480
gecaaggaac igeecialti igaaggigat tietggeeca aigigtiaga agagageatt
aaggaactag aacaagaaga agaggagagg aaaaaggaag agagcactgc agccagtgaa
                                                                   3540
accactgagg gcagtcaggg cgacagcaag aatgcc
                                                                    3576
```

<211> 4050

<212> DNA

<213> Homo sapiens

```
60
gaactttata gaaaggctag gcaaaaaatg agacccagag atatggaaga aactggccaa
agaggggaaa glgggctatt tettititt tittettitt tettietgag acagagtete
                                                                     120
accetginge ceaggeigga gigeagigae agegaieing geleacigea ageicegeet
                                                                     180
eccgggitea egecaticit eigecteage etecigagea getgigacia eaggigecea
                                                                     240
ccaccatgcc tggctaattt ttttatattt ttattagaga cagggtttca ccatgttagc
                                                                     300
caggatggic icgaleteet gacetigtga teegeeegee tiggeeteec aaagtgeigg
                                                                    360
gattacagge caeegtacee ggeeaggeta tttttttgtt tigititigt ttactacigt
                                                                    420
allgcillic iciliticat attiatigag cacciactat atgccagica ciatgciaga
                                                                    480
tgcttlagta acatgaaggt ttcaaactaa gaaaagctca acaaagagcc ctttagaaag
                                                                    540
glaacagili teetgiglat tgggggagtg ggteteataa ggttgtatga tgagaagege
                                                                    600
aagtaattgt liigittitt titgagacaa tgictigcic tgicgatcag geiggagige
                                                                    660
```

agtggcgtca	tctcagctcg	ctacaacctc	cacctcctgg	gtttaagtga	ttctctggcc	720
tcagcctcct	gagtagctga	gattacaggc	acgcgccacc	acgcccggct	aatttttgta	780
tttttagtag	agacagggtt	tcactatgtt	ggtcaggctg	atcttgaact	cccggcctca	840
ggtgatctgc	ctgccttggc	ctcccaaact	gctgggatta	cgggcatgag	ccaccacgcc	900
cagcctggct	aaggttcatt	atccccatct	ttcagatgag	gaggcacaca	acteteccea	960
gateceaegt	cacagagaat	gcaggtctga	gtctgttcct	tgagctcagg	gtcttggcaa	1020
tggtgatgct	tiggagcigc	agaatcctct	gttggggatg	gggctgccct	gtgcattata	1080
gtacgtttag	caacatccct	agcttccacc	tactaaatgc	cactagcact	ccttcagatg	1140
agaccaccaa	aaatgtctcc	agatgttacc	acatgtcccc	tggggggtaa	aagtgcccc	1200
gttgagagca	ctggctcttt	tcttcacagt	cctgacctgg	cggcctgcac	aggccacttc	1260
tccgaagtgt	ttcaatgcac	tctctgccct	gggtaccttg	gacacagcac	cctggcccag	1320
agaggttggc	tgacttgcct	gagacgctgc	ttcctgggag	acgcagtagc	atctttcctt	1380
tctgttctgg	ttatctttct	tagttcttta	ccaccttata	ttccccatga	caggtgtgtt	1440
tatgtacaca	catctgcctc	actccactca	gctccctgtc	aggtttcctg	ccagtctgtc	1500
cctcttcctt	caggctcagc	tacgtcctgc	acagacagta	ccactgcaca	tacctgtgtg	1560
tgcccagcgg	tggacccacc	tccaaaagca	gccagtgctg	acagcagaga	gccttccaca	1620
ctcaagtcag	gccaagcagg	aatcgctacc	tgcctgtcat	gaccacattc	tcagtgaaca	1680
ttgacaaagc	ccccttagca	gctaattagc	cctgccgtgc	gctagggatg	caatttctca	1740
tctggcagtg	cgccacactc	ctgcctccct	gcccaaagga	cgtagtggct	gctgctgatc	1800
gtctgcactg	ctgttccagg	ggcaggaggt	ttgctgcaaa	tcaggtaccc	ccagctcagt	1860
gagcagaacc	agtccaaggt	tgagtgagga	gaagggcaag	aagggcaggc	acagccgtga	1920
gtatgttctg	gggctaagta	accatgaggt	cagcccagag	accttgcaca	gttaggcagg	1980
cctggacttc	tegecettee	ccttgcagct	tctgctctcc	cagctaggga	ctgaggaaag	2040
ccctgcttct	agatgccatg	tgctgctgcc	tggcacgata	ggtacccatc	tgtccttggg	2100
gttcctgagc	ctggagagcg	ggctttgtga	gcactggtgc	ctcacctgcc	tggctcagct	2160
ctgcagccac	aatatatgct	taatacctat	tigitaaatg	attgaagact	tgactgccat	2220
teagtacaga	gaattagcca	ggtgaataaa	caggatgtgt	catagaggtt	ctagaattga	2280
teatgaceet	ttctgtctca	ttcctgactt	ctaataccgt	atatgccaaa	tggggttctg	2340
ctgtgattta	atttcttaag	gactgggttt	atcaaaagtc	cctcctgatc	taatcctttc	2400
ctctaggaag	getteteett	tcttcatctg	tectaagtge	atggtcttca	tctcctgggt	2460
ggtccagact	aggtggcact	gggcctgcag	gcctctagct	gctcaaggat	ggccctgtct	2520
gcatgettee	tttcaaaaagc	tagcatagaa	aggagggccc	aaggtgagga	aatttgtcca	2580
aagtcaccca	atgagtegea	ggaagggcta	gaatctggtt	atctggaccg	tectagagea	2640
cttlcacagt	gacagccggc	tggaatcaag	ttttcattta	gaaaaatggc	tagaagttag	2700
ggcattgcct	gcagccactg	aaaagcagct	ttaggagcag	atgtccacgt	aatagaagga	2760
gatgggctag	ggcctgccac	ggaagccagc	aagcgcgtgg	gagctggggg	aggaaaggag	2820

caaaaggcaa	gaacaggcag	tatgtccgcg	gtgcccacag	tgctgtgggt	acaagcaagg	2880
ggaaaagagc	ccatggtgtg	cagaaaacca	tgcgtcatga	ttcttatttc	ctgctcgcag	2940
ctttgactct	ctgcctcatc	tcttcctgga	agtgtcttgg	aagttaggcg	actgcacagg	3000
gaaaggttcg	ctgcagtgct	tgcaggcctg	cacccattta	ttcatccggt	ggatatttgc	3060
tgggtgcccg	gcctggggat	ccatggtgag	cgaggaaggc	atggtattga	agtggtatgc	3120
ctgcatgacc	ttggcggggg	cgcatggcat	agagaggaca	ggcttcagaa	caggcaggca	3180
agggctgaaa	tcctatctct	gccaccgaac	agctaatgac	cccagcaage	aatttcacat	3240
ccccgaactt	tcctgtttcc	tcatgtgtca	aatggggatg	atctcgagac	gactctccag	3300
agtaaccacg	tgaagcacct	agcacagggg	ctgacgcaaa	cagctgggca	tcggaggagc	3360
ctccagggtt	gtgacctcca	gtggcttatt	ttccttttgg	gatcttctct	cctagatcct	3420
cccctttaat	tccctgtgaa	atttaccact	ttcatattga	atcgttggca	cacagggcta	3480
actgcttgtt	cacctgaagg	aagctacaga	gttcaggttt	ctttttctt	tctttctttc	3540
ttttttgctt	ttttaagatg	atcttgctcc	gtcacccagg	ctggagtgca	gtggcgtaat	3600
catggcttcc	tgcagcctca	aactcctggg	ctcaatgagt	tccttgagat	cttccatcct	3660
cagcttccca	agtagctagt	agtagtagtg	gctlgcacca	acgctcctgc	cctaattttc	3720
aatattttt	tgtagagata	ggatctcact	gtgttaccca	ggctggactt	gaactcctgg	3780
cctcaggcga	tccttccgcc	ttggcctccc	aaagtgttgg	gattacaggc	attagctacc	3840
acacctggcc	aaggcccagg	tttcgacaga	aagggagaga	aaacctgcca	gagatgccat	3900
ttcggagcca	ctctgcttgg	cagggacctg	tgttcccctc	atgcaggttc	atccttagag	3960
ggctgcggtc	ttatctggtt	gtgcaaaagt	cccacaacct	ttctgggttg	atagtttgtg	4020
gtgaaataaa	caattttagt	ttgtttggag				4050

<211> 4907

<212> DNA

<213> Homo sapiens

ctttttgaga	cccttccctt	ggacagcatt	ggacagggtg	aggttctggc	ccatgggagt	60
ccaagcagag	aagaaggaac	tgattctgct	gggcaggccc	agggcatagg	gtccccagtg	120
tatgccatgc	aggacagcaa	gggccgcctc	catgccctga	cctctgttag	cagagagcag	180
atagtcggag	gtgatgtgca	gggctacagg	tggatgtttg	agacacagcc	cctagaccag	240
ctcggccgaa	gccccagtac	catcgacgtg	gtgcggggca	teacceggea	ggaagtggtg	300
gctggggacg	ttggcacagc	teggtggett	tttgagaccc	agcccctgga	gatgatccac	360

```
420
caacgggagc agcaggaacg acagaaagaa gaagggaaga gtcagggaga cccccagcct
                                                                     480
gaggeacccc caaagggega tgtgeagacc atceggtggt tgttegagac ttgeecaatg
agtgagttgg ccgaaaagca ggggtcagag gtcacagatc ccacagccaa ggctgaggca
                                                                     540
cagtectgea cetggatgtt caageeccaa cetgtggaca ggecagtggg etecagggag
                                                                     600
cagcaccige aggitageca ggiccegget ggggaaagac agacagacag acacgictit
                                                                     660
                                                                     720
gagaccgagc ctcttcaggc ctcaggccgt ccctgtggaa gacggcctgt gagatactgc
agccgcglgg agatcccttc agggcagglg tctcglcaga aagaggtttt tcaggccctg
                                                                     780
gaggcaggca agaaggaaga acaggagccc cgggtaatcg ctgggtccat ccccgcgggt
                                                                     840
                                                                     900
tetgtecaca agtteacttg getttttgag aattgteeca tgggeteect ggeagetgag
agcatccaag ggggcaacct cctggaagag cagcccatga gcccctcagg caacaggatg
                                                                     960
                                                                    1020
caagagagcc aggagactgc agctgagggg accetgegga ctetgeatgc cacacetggc
atectgeace atggaggeat ceteatggag geeegaggge eaggggaget etgtettgee
                                                                    1080
aagtatgtgc tetegggeac agggeagggg caccettata tacgaaagga ggagetggtg
                                                                    1140
teaggtgaac tteecaggat catetgecaa gleetgegee ggeeagatgt ggaccageag
                                                                    1200
                                                                    1260
gggctgctgg tgcaggaaga cccaactggc cagctccaac tcaagccgct gaggctgcca
actocaggoa goagtgggaa tattgaagac atggaccotg agotocagoa gotgotggot
                                                                    1320
tgcggtcttg ggacctccgt ggcaaggact gggctggtga tgcaggagac agagcagggc
                                                                    1380
etggtegeae tgaetgeeta etetetgeag eeceggetaa etageaagge etetgagagg
                                                                    1440
agcagcgtgc agctgttggc cagctgcata gataaaggag acctgagtgg cctgcacagt
                                                                    1500
ctgcggtggg agccccggc tgacccgagt ccagtgccag ccagcgaggg ggcccagagc
                                                                    1560
                                                                    1620
etgeacceaa etgagageat catecatgit ecceeactgg acceeageat ggggatgggg
catetgagag ceteagggge caecectige cetecteagg ceatiggaaa ggeagteeet
                                                                    1680
                                                                    1740
ctggctgggg aagctgcagc accagcccaa ttgcaaaaca cagaaaagca ggaagacagt
cactetggac agaaagggat ggcagtettg ggaaagtcag aaggagccac gactacccct
                                                                    1800
                                                                    1860
ceggggeetg gggeeceaga ceteetggee geeatgeaga gtetgeggat ggeaacaget
                                                                    1920
gaageeeaga geetgeacea geaagttetg aacaageaca ageagggeee caceecaaca
                                                                    1980
gecacticea accecateca ggacggicti eggaaageig gggetaceca aagcaacata
aggcctgggg gtggaagtga teeeeggate ceageageee eeagaaaggt eagtagggaa
                                                                    2040
                                                                    2100
gagcaagcac tacccagagg gctgcctggg gggtgggtga caattcagga tggcatctac
accectcate cegtgaggae ctttgaccea cetgggggtg tecagettte teagagggaa
                                                                    2160
                                                                    2220
ceccagicaa ggeacaggga gactgeecte teagleeagg eteccegece aciecaggga
ggcccaggtc agagtactgg gccagggcgg gaggagcctg ggggctgcac acagatggcc
                                                                    2280
                                                                    2340
tgggggccac cagggaaggc gatggcagaa gtctgcccag ggggcctcca agctgcagag
accaecetga agaetgeece tetaggeege cacattetgg cetetgggee ecaagetgea
                                                                    2400
                                                                    2460
ggtgccagcc cgcacccca taatgccttt gttcctcctc ctcctactct cccagctgct
gtgacaggac ctgactttcc agctggagcc caccgtgctg aggactccat ccagcaagcc
                                                                    2520
```

tctgagcccc	tgaaggaccc	ccttcttcac	teccacagea	gccctgctgg	ccagagaacc	2580
cctggagggt	cacagacaaa	gaccccaaaa	ctggacccca	ccatgccccc	aaagaagaag	2640
ccgcagctgc	cccctaaacc	tgcacaccta	acccagagcc	accetectea	gaggctgccc	2700
aagcccttgc	ctctatctcc	cagcttttcc	tcggaggtgg	ggcaaagaga	acaccaacga	2760
ggtgagagag	atacagccat	ccctcagcca	gccaaggttc	ccactactgt	agaccagggc	2820
cacatacctc	tggccagatg	tcccagtgga	catagccagc	ccagcttaca	acatggcctc	2880
agcaccacgg	ccccaggcc	caccaagaat	caggetacag	gcagcaatgc	ccagagetet	2940
gagcccccca	agctcaatgc	cctcaaccat	gatcccacct	caccacagtg	gggccccggc	3000
ccctcaggag	agcagcccat	ggaaggttcc	caccaagggg	cccctgagag	ccctgacagt	3060
ctgcaaagaa	accagaaaga	gctccagggc	ctcctgaacc	aggtgcaagc	cctggagaag	3120
gaggccgcaa	gcagtgtgga	cgtgcaggcc	ctgcggaggc	tctttgaggc	cgtgccccag	3180
ctgggagggg	ctgctcctca	ggctcctgct	gcccaccaaa	agcccgaggc	ctcagtggag	3240
caggcctttg	gggagctgac	acgggtcagc	acggaagttg	ctcaactgaa	ggaacagacc	3300
ttggcaaggc	tgctggacat	tgaagaggct	gtgcacaagg	cactcagctc	catgictage	3360
ctccagcctg	aggccagtgc	cagaggccat	ttccagggac	ctccaaaaga	ccacagtgcc	3420
cacaagatca	gtgtcacagt	cagcagtagc	gccaggccca	gtggctcagg	ccaggaggtc	3480
ggaggtcaaa	ctgcagtcaa	gaaccaagcc	aaggttgaat	gccacactga	ggcccagagt	3540
caagtcaaga	tcagaaatca	cacagaggcc	agaggtcaca	cagcctcaac	tgccccttcc	3600
accaggaggc	aggagacatc	aagagagtat	ttgtgccctc	ctcgggtttt	accitccagc	3660
cgagattctc	cctcctccc	aacatttatc	tccatccagt	cggccacaag	gaagcctcta	3720
gagactccca	gctttaaggg	caaccctgat	gtctcagtga	aaagcacaca	actggctcag	3780
gacataggcc	aggccctgct	ccaccagaaa	ggtgtccaag	acaaaactgg	gaagaaggac	3840
atcacccagt	gctctgtgca	acctgaacct	gccctccct	cagccagtcc	cctgcccaga	3900
gggtggcaaa	agagtgttct	ggagctacag	acggggccag	ggageteaca	acactatgga	3960
gccatgagaa	ccgtgactga	acagtatgag	gaggtggacc	agtttgggaa	cacagteete	4020
atgtcttcca	ccacagtcac	cgagcaggca	gagccaccca	ggaacccagg	ctcccacctc	4080
gggctccacg	cctcccctt	gctgaggcag	ttcctgcaca	gcccagctgg	gttcagcagt	4140
gacctgacag	aagctgagac	ggtgcaggtg	tcctgcagct	acteceagee	agctgcccag	4200
tgaggcccac	cgcctcccac	cacacctgcc	accigitect	ggcctccact	gccccaggac	4260
tgaagtgggt	acctgcctcc	tgtacactgg	agcaaggacc	aagaggaaat	ggcatcttca	4320
gaggattact	gtgggccatt	tccctttcgc	agtictica	ataggeceag	ttcttccaaa	4380
tggaaaaaga	aaggtctgga	agaggcccac	agagttgcac	aggcgtgggg	gtaggatggg	4440
ggctcccagc	tgcttgtgga	ggatgtaata	tatacagaca	cacacatgtt	tttcacacag	4500
gcctggccca	cgcatcgaca	tgtgtgaatt	tgcacaccac	tgcctgaatt	ggagececee	4560
agagtgtccc	tctacccaga	gtttttattt	ctttaattag	tctgagtgtt	cccagccatc	4620
tgctccttaa	tccctggaga	ggaacagagc	caactggaca	cagcgttggt	ctctgtttgg	4680

aatcactgtg	${\tt aggtctccag}$	aaggacctgg	ccgccagccc	cttcatcacc	atctccatca	4740
ttcagctggt	catctggtgg	cccaaaggtc	acccaaagag	tcagcaatca	gcatgtccct	4800
agaagccaaa	tgcactgcct	ttctctgtcc	ccatgactgt	ccccactct	gcaccccaaa	4860
tgggaagcat	acggtctgaa	taaatccaag	ttttattctc	tactctg		4907

<211> 4964

<212> DNA

<213> Homo sapiens

<400> 2010

60 agegggegee getageeage ggaagatgge ggagggegga ggeeetgage eeggegagea ggagaggagg tollcogggc ogcggcotco gagogoggg gatttgcagt tggcottggc 120 agaattgtat gaagatgaag tgaagtgcaa atcttccaag tctaatagac ctaaagccac 180 agtetteaag ageceaegga caccacetea aeggttitae teaagtgaae atgaalaeag 240 tggattaaat atagttcgac cttcaactgg gaaaattgtg aatgaacttt tcaaagaggc 300 aagggaacat ggggctgtcc ctctgaatga agccacaaga gcttcaggtg atgataaatc 360 taagtcattt acaggtggag gatacagatt gggtagttct ttttgtaagc ggtctgaata 420 480 tatctatgga gaaaatcagc tgcaagatgt tcagattttg cttaaactgt ggagcaatgg 540 tttcagttta gatgatgag aattgagacc ttacaatgaa ccaacaaatg ctcaatttct ggagtctgtt aagagaggag agattcccct ggagcttcag cgccttgttc atggtggcca 600 agtgaattig galaiggagg aicaicagga icaagaalac alaaaaccia galigaggii 660 caaggotttt agtggagaag ggcaaaaact tggaagcott acaccigaaa tagloagtac 720 780 accticctet ccagaagagg aggataaate aatacttaat geagtigite tialigatga ttcagtgcca acaacaaaaa ttcaaatcag gttagcagat gggagtcgtt tgatacaaag 840 attcaatagt acacacagga teetggatgt eeggaacttt attgtacagt etegteetga 900 attigegget etigaetiia iteitgigae iteallieeg aalaaagage taacagaiga 960 aageetgaca etgetagaag cagatattet taacactgig tiactecage aactaaaata 1020 atatigitcc igiccatgca glagcaigig ggaalagaig aigigccgia ilaalaagga 1080 caatactica gcallaaaaa cagccaaall attittatta tiillacaga taaatiligg 1140 tittaitgit alicigicii ccaaicigaa tatagacaaa iiiggaltag gaalagacci 1200 tgagataagt algittgagi iittagilga aggaciggci talgitgala gillilggal 1260 ttclaggcaa algagitgit acaigcitag igitaaigia acaacaiiig iiigcagaga 1320 aaaatgaaca aaaccccttt tigalaaatg catiigglaa aatiigcact aaagtiictt 1380 gatgcagcat tgaccaacag ccattaagaa atctttigat caaataagtt gaaaattigi 1440

```
1500
ctataatata tactgaaacg tgtcttttga ttttgaaatt gtttgatcat acaataatta
tttctcctat taagatttta cacatccttt ttacttactg atttagatat attactagta
                                                                    1560
                                                                    1620
tcagaaacta cagttttgcc ttgtatttta cagaattatg actgttgtga acttaaacag
aaacacataa aggtcagcaa ttctttttt ttttttttt gatatggagt tttgctcttg
                                                                    1680
ttgcccaggc tggagtgcaa tggcataatt tctgctcacc gcaacctccg cctcccaggt
                                                                    1740
tcaaaagatt ctcctgcctt agcctccaa gtagctggga ttacaggcat gcgccaccat
                                                                    1800
gcctggctaa tttttgtact tttgctagag acagcgtttc tctgtgttga tcaggctggt
                                                                    1860
ctcgaactcc gaacctcagg tgatccaccc acctcagcct cccaaagtgc tgggattaca
                                                                    1920
                                                                    1980
ggcatgagcc accacgccca gcctaaaggt cagcagttct taagaagata tggtaaacag
                                                                    2040
caacaatatt ttaaaatcaa gtaattacag ttcctcccag agcttgcgtt gatcacattc
                                                                    2100
atttatteat teaacacatt tttetaggaa acteactgta tacactaaac actattetgt
gtgctcaacc tagaatgtct tctccagaac aagactagtg tagaaataca ggaatgtaaa
                                                                    2160
                                                                    2220
ttctgtcaga cggactagat ctaaagaatt accagcataa atgtttgcat ttctgctgaa
                                                                    2280
gecagaaget ttteettett eetagacaee attteateet taattattae ttetggttag
ttttccattg ccaccataac aagttacaaa atgtggctta aaatagcaca aatttattat
                                                                    2340
cttcacaatt ctgtaggtta ggagtccagg ttaagagttt cgcggtgcca agatcaattt
                                                                    2400
gttggcaggg ttgcattctg ttaggaggct ctacaggaga atcatttcct tgtcattcca
                                                                    2460
                                                                    2520
cettetacag gacatectea tteettgget tgtgacetee ttetteeate ttaaaaacea
                                                                    2580
gtgctgtttc atctctatga cccttctgtt accacatctc tctgacacca gtgtggagag
                                                                    2640
gttctctgca ggactcatga ttaaatgagg cccaccggat atccaatcta ggcttatctc
                                                                    2700
ctigictiga aatccatagt aaccttaatt acatctgcaa aatctciitt accatctaag
gtlacataca ggtttggaga tlaggacatt aacattttac atggaacatt attcttgcct
                                                                    2760
                                                                    2820
actacagtic ccacccacce eccgetecae teetgigita aagatteaga ticatcacaa
                                                                    2880
ataaatttac atcactcata ggtgctcaaa agtcacaatc cattattaca gcatcaactc
                                                                    2940
taaatccaaa atettatetg agteteacea acteaaaagt eteaaatete acattgaage
                                                                    3000
catclaaatt aagtttggga gaggatctgt gigtgatiic tgggacataa iiccaactgt
                                                                    3060
gcactigiga acctagaaaa caagitatci giicccaagi algatggcai gacaggcaga
                                                                    3120
caalaalagt tacacacgtt cetgilcaaa aagcagaaac agalggaaaa aggagccatc
agcaccaate aatttacaaa accagegagg caccettett taagtticaa ggeetgggag
                                                                    3180
taatetteag eteaetgetg ttetetggge tigitgaetg teleagagte atetitaett
                                                                    3240
ttlcacaaaa ggtagcacac gtltgcagct gagtatcaac ttalcagitt gttcttcttt
                                                                    3300
talattetet aaagetitet gitaaaaaig giggigelte igetgetata aegitgicaa
                                                                    3360
                                                                    3420
gaaactigig ggictittac ataigicaca gggaigeact cattiagata ggaggeteel
cacgtatett teetggaaaa teetgtetet gtttttgget tiltetgaaa tagetgagag
                                                                    3480
                                                                    3540
gatelatgat teacaccett aatatettea aagagtettg tgtgtgacet gatatteaga
                                                                    3600
cclittgatg itlctgaagt attagcaaaa ggliatacag ccatatciic atcactiici
```

```
ctagagtaaa ggctgtcctg acggtgaatc ttagttttag tggcttttgc catttgaata
                                                                    3660
ggccgcgaat ttcccaaatc atcaagtcct ggtttcttta tatttaacag gtcttcctc
                                                                    3720
aatctacctc tttccacatt ttactataat cagcaagaag acagcaggct gtaccttcca
                                                                    3780
cagctigctt ggaaatatcc tcagctaaat attgaagtca tcacttaaaa gttctgcttt
                                                                    3840
acacataacg gcaggacaca actcagctta gcttttcgcc actatgtaac aaggactcct
                                                                    3900
tlectecact tetecagtaa catatteete attttttace aacagtetat teatgatgat
                                                                    3960
ttagatatte tatggeaate gaggtattet etattatget eetttettea aggeegeeet
                                                                    4020
agcallaaca ttccatattt ctactaacag tctgtttaag gcagtttagc ticttttctg
                                                                   4080
gcatgctcct cagaattctt ccagcctcca cctactgccc aattccagag ccacttttct
                                                                    4140
acttttaggt atttgttaca gcagcacctc aagtacctag aaaactcttt tatgcctgct
                                                                   4200
tetetgecag atgaettgaa tatggtaeta gatttggaat teaeetttet eeagggteae
                                                                    4260
tgtttatttc aaagaggtga atttacctgt gctagggttt tcacactggg agtgctacca
                                                                    4320
gaactaccac aggatgaaag tggtgagccc accactgcag agaagttttc tcagtgccgt
                                                                   4380
aatatagagg aattotoaaa ataagooota otoottitoa ottaotgaaa acaactigga
                                                                   4440
taatgtgtaa cagccagccc catttcaaaa agattaccag gggtaaaaca actttttcat
                                                                   4500
gggtcaaaat catcttccga agaaaatgat ttcttaaaag aattgaacat tgtaaatcaa
                                                                   4560
agggcattgt cctgttttgg attaacaaaa caggaaaaat aaccaatcct tgtaaaatta
                                                                   4620
tttgaaattt tcttgttttt atcagttgag tgcctataga tgcacataca aaaacaactg
                                                                   4680
ccatttttgt atataatagt cttccaagat agagatttac attaggagag aattaaacat
                                                                   4740
ccaggaggga tgaacagtat ttcatgtgtg ctatgtagtg ttttgcttca ttgagagtca
                                                                   4800
tittcatgaa tiattittac tacigcagic atcitaaati talaatcatc tcaaaaaaga
                                                                   4860
tgtcacaatg aacagacaac catctgtgag gtcagtcatt ttgcatgatg tatgtaatca
                                                                   4920
aaaagttiga aatgicigci tactaataaa gaatgittic actg
                                                                   4964
```

<211> 3825

<212> DNA

<213> Homo sapiens

ctttcctttt	cgcctctcct	cgttctctcc	ctcgcctttc	ctttcctttc	tcttcctctc	60
ttcctcgctc	ctcggtctcg	gcgctctccc	agcttttctt	ctcctggctc	ctggttcccc	120
gctacgccac	cagtccactc	acctetetee	ttgccctact	ccetecgeta	ctccctgacg	180
cccctgcag	ccccagccc	ccctgcaggc	cccagcccca	gtaagtttgg	agaggggaac	240
aaatgctgag	cctaggtagg	gaccaccttg	gggaggaagc	caaaatcaca	ctgctcaccc	300

```
gagageeeet geeeeget ggeaegeeee egeetggagt geaetegtgg eeeegggege
                                                                     360
                                                                     420
tgtcaggtac ccgaattggg gctgccaccg tgtcggaggc gaggcgagga agggagctgg
aataacaaag gtggcagctg agcatccctg gagagggtgg gtggtatgaa agcacttcca
                                                                     480
gacctctagg gacaccaggg agtcatggtc ccagcacatt gctgtgtgat tgagccctc
                                                                     540
ctcagcctgt gggtggccta agttcacagg gaggtaatgg ggtagattgg atacctctgg
                                                                     600
ggtcttggaa gaagctatga cttatttact gtctactatg tgatgggaag ataagaccca
                                                                     660
gaaaacagaa aggacatgtt taaggccatg cagcaagtta gtgcctgacc tgaatattga
                                                                     720
agtgaggece tactaceate agecatggga accatggetg gatgggtece aageaatgaa
                                                                     780
gaccttctgg gtgtctaggg gagaggtttg ggccctcctc catgtgcgtg tgtgtgtgcg
                                                                     840
cgtgcaagtg tgtgtgtctg ggaagccaga agattacact cttctttcta ggccttctag
                                                                     900
cccttgctgg aaggcctgta gtgagtggat ggcctgcctt accctctgca catcccgccc
                                                                     960
tgtttattga gatttccatc cagcctgaac tcctgtgggg aggtgttatc ttctggacca
                                                                    1020
gagecetate tgeealgaag ceatigiggi gicaeagggg eticigagag ateceagget
                                                                    1080
ggagacggaa agcagaagat ttgaagtggt gggaggcagg ggctggtgtc ataacacact
                                                                    1140
ttecaeccet gggetgggag gggeacteec teetgetgaa eteteecagg ecagtgacet
                                                                    1200
catettgete etgtgettgt titceaaagg gigtiglaag tigacigiet getitettee
                                                                    1260
acaacactca aagtgtggcc tgtggagcaa cagcttcagc cacagctggg agctggttag
                                                                    1320
aagtgcaaca totcaggcoo caccotagaa cattaacato totggaggta ggaccoagga
                                                                    1380
atetgtttea caagteetet tetgatgett agaaaagttt aaacateact getttaetet
                                                                    1440
atttctcgac aaaaagatga cattcagttt ggctagaatt aaaaggggtg ggtgtttcct
                                                                    1500
ggcaggtttt agaaacctat ttaaatggtt ccattgtcca ttcatccatc catcaaccca
                                                                    1560
tecaaceate cagecageea tecaecetet titeatteaa cagacattea getgeacteg
                                                                    1620
ggagttgaaa ggggaagget egggaeeetg ggeteeteea gettgeegtg agacaceaet
                                                                    1680
gtgtggcaga agaggtggcc tctgtcccct ttatcctcca agtgtacctg tggtcttcag
                                                                    1740
geeggteact tgettgaate tgagtgtgt tetetgatet ataateetaa aaaagetace
                                                                    1800
taatgcaggt gtccaagagg gaaaggggaa ggaattgcat gcacttggtg tctattgtgt
                                                                    1860
gccaggtgtg ttcacacgtg ttatgacctt ccactcttcc agctgccctt catactggat
                                                                    1920
agcattatto ttattttaca gagaagaata ttgaggatca aagagaccaa gactgcaagc
                                                                    1980
gtaaaaacta agattggaac caaagccagt tettetegat eccagggtet gegeeettet
                                                                    2040
tetgtteeat gitteatigi tetiggigga eeiggggate aatagetaga agitaaagga
                                                                    2100
caaactgatt tgggaagtgc ttccagtgct gtcttgagtg atgtctagag attagcagac
                                                                    2160
tggctgtgaa gtggtgagct gcccatcact ggaaccgtgc aagcagagac tggtcatggt
                                                                   2220
gatcacggig tiggiccigi gigagigiga tgiggggaag aatigagacc agaigaccii
                                                                   2280
tgagggcctt ctgctgtctg aggcgggcct gcttgggcct gctcccaggt cagtgccca
                                                                   2340
tggatggage etetgaagee agetgeteat tatetgtgga teetetgegg ggacaetgee
                                                                   2400
ageteceaaa caggaaacat gtecagaaat etgtaattag agetgggage cacaggeetg
                                                                   2460
```

agaggtgcct	gctgcagctt	caagtgcaga	cacgccaccc	tggttaagtc	cctgggagag	2520
aacccagtga	gtcaggccct	cagatettet	ccctgcctgt	ggcacccccg	ccccacccca	2580
ttccccttgg	aaggaaacct	gctttggcca	ggaacctact	gggtgaatgg	gtttcatata	2640
cattctctcc	tctgttcttc	cccagaaccg	tgggagagag	gaaacatctg	ccatgatgca	2700
ggcaaggaat	gcaaagctcc	cagacatcat	gtggctcact	caaggtcacc	ctactatggc	2760
ccttgccttt	ctgagtgcct	ggtttgacct	cttgatccct	ccaggggaga	acgtcacagt	2820
caaaggaggg	gtgcaagagg	ccagtggcac	acagagaggt	ctgtgtgggc	ctgagtggct	2880
cctgggtctt	ccctgactga	ccataacgcc	tttcagcctt	tctgaatctg	ccatgaaggg	2940
acgggtcctt	gcagtgttcc	tctgccaggc	tgcctggcaa	cccatggcaa	ttgtggtggt	3000
gttaaaacat	ggccacaggc	caggcacggt	ggctcatgcc	tgtaattcca	gcactttgca	3060
tagggtatgg	cagaagagac	cctaagtgag	taaagaccat	gcccctgcaa	attatacttt	3120
gtttgctgga	acattcactc	ttggagccct	gagccaccat	gtaaagaagt	aggaagattc	3180
actgtcctga	agctgccatg	ttgtgaggaa	gcccaagcca	catggagggg	ccatgtctgg	3240
gtgctccggt	caacagtccc	agctgagctt	agccatctaa	catccccagc	tattttagtt	3300
tttcctgaaa	tcccagaaat	catggaatgg	agacaaatct	ctcctgctgt	gctctgtctg	3360
aactgctgac	ccacagaatc	tgggcacata	ataaaattat	tttgtgccat	taggtatata	3420
gttgatttgt	tatgcagcca	tagataacca	ggacagctat	gccagctatg	aagtgccatg	3480
cagtcatctc	gggggtccca	ctcacaacat	ctccccatac	tcctaggaag	ctggctgggc	3540
tcaactctaa	gtgcaaagca	ttgtgcaaag	ggaagggcat	gaaactgggg	ggccctgcat	3600
ctcctggggg	ttagagtact	gaacttcctc	cacccactgc	cttctcagag	atgagcaccc	3660
tacatctgga	tctgcctcag	gccctcttgt	atatgactaa	gaatattggc	ttggtgtggt	3720
ggctcatgcc	tgtgatcccg	gtactttggg	agactgaggc	gggaggatcc	ttgagcccag	3780
gagtttgaga	ccagcctggg	caacacaaca	agaccctatc	tctac		3825

<211> 3483

<212> DNA

<213> Homo sapiens

ttgaaaatat	tttcatgaga	atttaaactg	acaaaaaatc	tagaagtttc	ttcttgcctg	60
agaccccccc	tcccagaaat	aatctctgct	atcagggtgt	gttctttcaa	gcctatttct	120
atgtatttgc	tcatatatag	aaatattict	agaatgatat	aggcttctgt	gttttattat	180
ctaaatcagt	cattettaac	caggggtgat	tttgtacccc	ctcctcctag	gagatacttg	240
gcaatgtctg	gagatatttt	tggttgtcac	acatagaggg	ggtgctactg	ccatctagta	300

ggtagagaga	ccaaggatgt	tgctaacatc	ctatagggca	caggacagcc	cccacaataa	360
agaatcaacg	tggcctaaaa	catcagtagt	gctggctggg	ctcacgcctg	taatcccagc	420
acttttggag	gccaaggtgg	gcggatcacc	tgaggtcggg	agttcaagac	cagcctgtcc	480
aacacggaga	aaccccatct	ctactgaaaa	tacaaaagta	gccgggcgtg	gtggcgcatg	540
tctgtaatcc	cagctactca	ggaggctgag	gcaggagaat	cacttgaagc	cgggagggag	600
gtggaggttg	cggtgagccg	agattgtacc	actgcactcc	agcctgggca	acaagagtga	660
aactctgtct	gaaaaaaaaa	aaaaaaaatt	atcagtagtg	ctgagaaacc	ctggtctaag	720
tggtggtgta	tggtatacat	tgttagacaa	tttcttttat	acaatgtttc	tgggtcagtc	780
tatttagatc	aactgatcgt	tttgcttact	gccaagtttt	ccatactacg	catagcaggt	840
agtcgagttc	accattcccc	atttagtgga	catctagacg	gctgctcgtt	tttatcattg	900
cagcattctt	tgcacacatc	cttggatatg	agcagacatg	aaaatgtttt	tctagggttg	960
acactgagca	gtaaaagtgc	tgggttgaag	ggtttccagc	ttgcatttgt	acctggcctt	1020
ctacagggga	cagggggcta	tttagatggt	cccctgccaa	ccccagtgga	caaccctagg	1080
gtggggctgg	aggtggggct	gaggctgagt	cttcctcccc	ttcctccctg	cccaggggtc	1140
cacattcagt	cgtcccagac	tgtggagtcg	agtggtttgt	acaccttgca	gagtattctg	1200
aaggcacagc	tggttaaaga	agacaaagat	gcccagtttt	actgtgagct	caactaccgg	1260
ctgcccagtg	ggaaccacat	gaaggagtcc	agggaagtca	ccgtccctgt	tttctacccg	1320
acagaaaaag	tgtggctgga	agtggagccc	gtgggaatgc	tgaaggaagg	ggaccgcgtg	1380
gaaatcaggt	gtttggctga	tggcaaccct	ccaccacact	tcagcatcag	caagcagaac	1440
cccagcacca	gggaggcaga	ggaagagaca	accaacgaca	acggggtcct	ggtgctggag	1500
cctgcccgga	aggaacacag	tgggcgctat	gaatgtcagg	gcctggactt	ggacaccatg	1560
atatcgctgc	tgagtgaacc	acaggaacta	ctggtgaact	atgtgtctga	cgtccgagtg	1620
agtcccgcag	cccctgagag	acaggaaggc	agcagcctca	ccctgacctg	tgaggcagag	1680
agtagccagg	acctcgagtt	ccagtggctg	agagaagaga	caggccaggt	gctggaaagg	1740
gggcctgtgc	ttcagttgca	tgacctgaaa	cgggaggcag	gaggcggcta	tcgctgcgtg	1800
gcgtctgtgc	ccagcatacc	cggcctgaac	cgcacacagc	tggtcaacgt	ggccattttt	1860
ggccccctt	ggatggcatt	caaggagagg	aaggtgtggg	t gaaagagaa	tatggtgttg	1920
aatctgtctt	gtgaagcgtc	agggcacccc	cggcccacca	tctcctggaa	cgtcaacggc	1980
acggcaagtg	aacaagacca	agatccacag	cgagtcctga	gcaccctgaa	tgtcctcgtg	2040
accccggagc	tgttggagac	aggtgttgaa	tgcacggcct	ccaacgacct	gggcaaaaaac	2100
accagcatcc	tcttcctgga	gctggtcaat	ttaaccaccc	tcacaccaga	ctccaacaca	2160
accactggcc	teageactic	cactgccagt	cctcatacca	gagccaacag	cacctccaca	2220
ggtaagccag	gcctggcaag	agaacagggc	tgtgccaggg	catcctttct	gccctgtccc	2280
tccccagaga	gccctgtcca	gaaaggtgag	tagcagcccc	atcttgtcgg	ccctggactg	2340
gctggggcaa	cgatggtgac	gaagtggcct	ggggcaggga	gtgacgagga	gtgtctttgt	2400

ggcgcagaga	gaaagctgcc	ggagccggag	agccggggcg	tggtcatcgt	ggctgtgatt	2460
gtgtgcatcc	tggtcctggc	ggtgctgggc	gctgtcctct	atttcctcta	taagaagggc	2520
aagctgccgt	gcaggcgctc	agggaagcag	gagatcacgc	tgccccgtc	tcgtaagagc	2580
gaacttgtag	ttgaagttaa	gtcagataag	ctcccagaag	agatgggcct	cctgcagggc	2640
agcagcggtg	acaagagggc	tccgggagac	cagggagaga	aatacatcga	tctgaggcat	2700
tagccccgaa	tcacttcagc	tcccttccct	gcctggacca	ttcccagctc	cctgctcact	2760
cttctctcag	ccaaagcctc	caaagggact	agagagaagc	ctcctgctcc	cctcgcctgc	2820
acaccccctt	tcagagggcc	actgggttag	gacctgagga	ccccacttgg	ccctgcaagg	2880
cccgcttttc	agggaccagt	ccaccaccat	ctccacgttg	agtgaagctc	atcccaagca	2940
aggagcccca	gtctcccgag	cgggctggct	tccaccatcc	aggtgcacca	ctgaagtgag	3000
gacacaccgg	agccaggcgc	ctgctcatgt	tgaagtgcgc	tgttcacacc	cgctccggag	3060
agcaccccag	cagcatccag	aagcagctgc	agtgttgctg	ccaccaccct	cctgtctgcc	3120
tcttcaaagt	ctcctgtgac	atttttctt	tggtcagaag	ccaggaactg	gtgtcattcc	3180
ttaaaagata	cgtgccgggg	ccaggtgtgg	tggctcacgc	ctgtaatccc	agcactttgg	3240
gaggccgagg	cgggcggatc	acaaagtcag	gacgagacca	tcctggctaa	cacggtgaaa	3300
ccctgtctct	actaaaaata	сааааааааа	ttagctaggc	gtagtggttg	gcacctatag	3360
teccagetae	teggaagget	gaagcaggag	aatggtatga	atccaggagg	tggagcttgc	3420
agtgagccga	gaccgtgcca	ctgcactcca	gcctgggcaa	cacagcgaga	ctccgtctcg	3480
agg						3483

<211> 4717

<212> DNA

<213> Homo sapiens

ttacttcaaa	cgggactcga	cccatgacca	cacctccaac	ctctctgccc	gagccctttt	60
ccggggaccc	aggccggttg	gcggggttcc	tgatgcagat	ggacagattc	atgatettee	120
aggcctcccg	cttcccgggt	gaggccgagc	gtgtggcctt	ccttgtgtct	cgactgactg	180
gggaggcgga	gaagtgggct	atcccccaca	tgcaacctga	cagccccttg	cgcaacaact	240
atcaggggtt	cctggcagag	ttgcggagaa	cctacaagtc	tccgctccgg	catgcgcggc	300
gcgcccaaat	caggaagact	tctgcctcta	atagggctgt	gcgagagagg	cagatgctct	360
gccgccagct	ggcctctgcg	ggcacggggc	cttgcccagt	gcatccagct	tccaacggga	420
ctagtccagc	gccagccctg	cctgcccgag	cacggaatet	ttaagaatcc	gccagcactt	480
ggtagcgtct	gcagccaccc	aggtagcata	cgctctttgc	tgtgtagaag	aaatgcccat	540

acgacagctt	tgccctgtt	tgaagacctc	ccttcttgcc	tctccagacg	tgttccccga	600
ggagatette	cttccgtcct	tcctggcgcc	ctggttgccc	accttgccgt	gcttcctctt	660
acgtgctagc	tttgtaccta	tcgctcactg	catgctcgcc	tccctcttgc	tggcatcccg	720
gcctgtttca	atgactaccg	ctctgctact	taggcacagg	gactccgccg	cacgctgacg	780
gaccacgagg	gctgacccct	tccagcctga	cttggttcat	ggaggctcct	actctgccct	840
ctccaagctc	ccctggcggc	tccccacctg	gttgcccagt	tcctattgat	gagctctgga	900
cagaaagatg	cccgtttggc	caggctggtg	gcttgatggg	tgtacctgga	gagggggtct	960
ggcttcctgc	ccaagatgcc	tcccagccct	gccagggccc	ggtgcagcgg	gcagggcctc	1020
atctgtgctg	tagtggtcga	gtggttgctg	caaggagcgt	agttctgcca	tgtctggggg	1080
ccaggttcca	ctctgcacat	gaatatgcag	tctgggaggc	cccactgctc	tcactgggaa	1140
ggaccaatgt	tgcacctctg	ttaatgcctg	acttcagctg	ctggtgttct	gatggagcca	1200
gaggcttggg	gaatctggaa	cttgcctgct	aaataaggtc	gtggtggact	ctcagccatt	1260
gggcaggtct	atcaggctgc	aggttcctac	acacccacgc	ctgagggtca	tagcaggcta	1320
agggtggata	ccagcgactc	cctttgctgc	ccaggatctc	catgggcagt	gccacagcgg	1380
ctgatgctca	gtcactcctg	cttctacccc	ctgtcactga	tggcgagcct	tgcccagctt	1440
gagacctgtt	cccatctcta	ttcaggtgcc	atgtggcctt	cactgcagcc	ctgcagccac	1500
ccacgcacca	tctgtgggtc	tccaaaggca	ccttgtagca	tgtactcccc	gtgcctgggc	1560
aatcagatgg	gctgcctttg	tccaagggaa	aacagactcc	cttcgggaaa	catccttaag	1620
cacttaaggc	cggggggggt	gtctgcctct	ggcaacccag	ccagggtctt	ggtggcattt	1680
gtaaaagcaa	agagctgtgg	actgccgtgg	tcctagtgtg	gtgacaatgc	agcactggca	1740
tgcatgtcct	ccttctgaag	gacctcatcc	ttcctcacag	ggggatgacc	aagaaatcat	1800
tttgtggctg	agtttggcca	cgccctttgg	actgtgctgt	tccgccatat	ttcaatgcca	1860
aatgaaccac	attgacatga	cctggaccat	agggcttcct	atcctgggct	cagctgcccc	1920
tgtctgaagg	gtcttggctt	gattgcagaa	ggacaacctc	cgcacccacc	taaagacatg	1980
tatatgtctt	gggatcccag	agattgggtc	cttgggcctg	gcttcttaag	agttttgatg	2040
atgctgggaa	aagtgactgc	gattctgaag	aaccgctgcc	ttgcaaggtc	aaggacattc	2100
agtggttgct	ggggtccgca	gactactgcc	acccactcac	catcaactct	gttagcccaa	2160
ttgccctgct	gaacaactgc	ctgaatacag	gctttaggtt	ccctggact	ccagccaagg	2220
ctgttcaggt	gggaccatgg	tgctctttaa	gcgtgatcgg	agggaagaca	cacagcaggg	2280
ccaccattcc	atgaatggga	ggtgtacaga	tcactttctc	tttgtgctca	gttctgttct	2340
gtctccagca	gctatattgg	taagactagt	acctgccagg	gagaggtgcc	cccaagtgaa	2400
ggggtacagt	ggcacctggg	aaaaggcacc	tggaaggttt	ccatgtggcc	cagcccagca	2460
	gtgggaactc	· ·				2520
	ccatgtctgt					2580
	acgctccctg					2640
ccctgtccac	${\tt aactgcagcc}$	gggccctctg	cctatgggca	cccaatccaa	gcagctgctc	2700

```
2760
cacctttgtt tggcatggtg atttgtattt tttctcttgg tgcttatgtg tgtgggcttg
                                                                    2820
ggacgagtgc tggtatgcac ttaggacctt cttgatagct ccctgcactt tggaacacgg
                                                                    2880
agcagatgag agagggtcgg ggcttgccct ccaccttgga cttggaagaa gcccacattg
                                                                    2940
gagaggtgag gaccccatgg tggctctagt ggaagatacg ttagtctcca gctaaggagg
atgaggcgca gccccagagg gagacctcag tgatagggga tcaggctaag aaagtggggg
                                                                    3000
                                                                    3060
aagggagatg ctttgtacat attttggggt tataatttct ctaaatttta ggagaacggg
                                                                    3120
tattgattga taaaagggac aggcagtagt gttcaacagt gcatgtgaag gaaagttctg
                                                                    3180
ttttccatgg ttttgacatt ctttggactg tattgtgact gctgtctggt ccacatggta
                                                                    3240
cccctttggt aagtaggett cagtgcatac cagggtatca ctggagatgg gagttagtga
                                                                    3300
aggggtgact ccctggccta gtatagtgtg accctgggac taacttaatg tcctaaagca
                                                                    3360
ttttggtgac ttctagggaa tagcaaagac ctatttcatt gtccccaggt aagtatgtga
tgagcaatga ggaggagtgg aaaacaaaac ccagaaagtg cggcaggacc agcctgacgc
                                                                    3420
                                                                    3480
acacgetect gttgtcatgg cagacageeg cettgggtgg gcaccaccet ggcagtteca
                                                                    3540
gcctgtaggg gagtgaaggg acatggctga gctgggcatg tgctgaggtt gacttaggga
                                                                    3600
acaagccctg ggattggaca aaagggccca tgctgcagcc actgactggg ggcagagctc
tgggtggaag agggaagaga tcctaatgga ggcgcctcca tctgcaacca cagttgtaag
                                                                    3660
gctcatggca cctctgcttg gaaagcactg gtttagggac ttagagaggt aggcacaagg
                                                                    3720
                                                                    3780
tgggtctcct gggtaaggga agcaagagca gactgttggg ccaacaggag aagctcccca
                                                                    3840
gagtagggga gaagattggg gtgtagggcc ttccacgtgg aacagacagc ccctgtgtct
                                                                    3900
ctgtctcttg gggacctgag tttgggtggg gtggcagttg gcacagcgca gatgcggtag
                                                                    3960
agalgggagg aaacccagct cctcacttcc gtgtgcctca tgcctttgca tacacaagca
ccaaacctac taggictict cattacccat giaaaccaca tgitagataa attitigcaa
                                                                    4020
                                                                    4080
gtagaggaaa gaaggaaata aaacatcaca ttttggtgtc tctcaggctt tccccccaa
ctatggtttc tttgcttttt gttttaacat agttttgttg ctgtcttctg taatgataca
                                                                    4140
                                                                    4200
gttttgtgca gctgttttca cttagcatat cgtgggcatc tccccttatg attactaaat
                                                                    4260
attitatitt ggagtggcig iglacicic cattgaciag atggaccatt gigccagiig
                                                                    4320
ccaatcacta atgctgttac taacttttca gttataaatt gatgaatatc tttgtgcaca
                                                                    4380
ggctgtttcc caatgtcaag ttattagggt agactccagg aggtgggatt cttcaactaa
                                                                    4440
agaatatgaa aacctttgag gcttttacta catattgaca aaatggittc cggaaatatt
tgtatcccct tacactgcca ccagcaagga taaacatgtc catcttgccc gtattgggaa
                                                                    4500
                                                                    4560
ttatcatcig gctaaatatt igctaattig ataatgaaaa aatagcatcg igtitcagti
ggcatttcac tgacttctag cacggttgaa catctttcat gtggagcgat tgtatttcct
                                                                    4620
                                                                    4680
cciligigga tigicagigi cciligcici atcitetggg gicagataaa titgiatgag
                                                                    4717
ctcggtatat attaaagata ttaacctggt gtgtgtc
```

```
<210> 2014
<211> 4112
<212> DNA
<213> Homo sapiens
```

```
60
attitatiga aggeettite igeatetati gggataatea igiagittiti gteatiggit
                                                                     120
cigittaigi gaiggattac gittaitgat tigcataigi tgaaccagcc tagcacccca
                                                                     180
gggatgaagc tgacttgatt gtggtggaca tgccttttga tgtgctctgg attcggtttg
                                                                     240
ccagtatggt attgaggata ttcacattga agttcatcag ggatattggc ctgaaatttt
                                                                     300
cttttttttg ttgtgtctct ggaggttttt ggtatcagga tgacactggc ctcataaaat
                                                                     360
gagtgatgga ggagtccctc tttttatatt gtttggaata gtttcagaag gaatggtacc
                                                                     420
ageteetett tgtacetetg gtagaatttg getgtgaate eatetggtee tgggettttt
tiggitgata ggcictiaat tacigciica atticagaac tigitatigg iciaticagg
                                                                     480
                                                                     540
gatttgactt ctttctggtt tagtcttggg agggtgtatg tgtccaggta tttatccatt
                                                                     600
tettetagat tittetagtgi attigeatag aegiattiat ageattetet gattgiaaae
                                                                     660
tgtatttctt tgggatcagt gatgatatcc cctttatcat tttttattgt gtctatttga
                                                                     720
ttcltctctc ttttcttctt cgttagtctg gctagtagtc tatctatttt gtgaatcttt
                                                                     780
tcaaaaaacc agctcctgga ttcgttgatt gtttttggtt ttccgtgtct ttatctcctt
                                                                     840
tggitctact ctgaicttag tiatticttg tcttctgcta gcilitgaat ttgtttgccc
                                                                     900
tigettetet igiteilite aligigatgg ggiatigalt tittatetti eetgettiet
cctgtgagcg cttagtgcta taaatttttc tctaaacact actttagctg tgtcctagag
                                                                     960
                                                                    1020
attetggtae attgtgtgtt eteattggtt teaaagaact tatttattte tgeettaatt
                                                                    1080
tcattattta cccagtagtc attcaggagc aggttgttca gttgccatgt agttgggcga
ttttcagtga gtttcttaat cttaacctct aatttgattg caccagggtc cgggagactg
                                                                    1140
                                                                    1200
ttatgattic igitetiitg eacitgetga ggagtgitti aciteeaati eigiggleaa
ttttagaata agtgtgatgi ggtgctgaga agaatgtata tictgitgat itggggtgga
                                                                    1260
gaglicigia gatgiciati agglicigett tgiccagage tgaglicaag teetgaatat
                                                                    1320
ccttgttaat tttctgtctc gttgatctgt ctaatattga cagtgggglg ttaaagtctc
                                                                    1380
                                                                    1440
ctactattaa tigggigga gictaagict ciligiaggi cictaagaac tigcilatga
                                                                    1500
attgggtgct tctgtatagg gtgcctatat atttagggta gttagctctt cttgttgcat
                                                                    1560
tgaacctttt accattatgt aatgeeette titgtettit tigateligg tiggittaaa
                                                                    1620
gicigilita icagaggela ggaligeagg altgeaacce etgettitti tititeitgg
                                                                    1680
tagatatice iccatilett tatitigage etaigtgigt ettigeatgi gagatgggte
                                                                    1740
tecegaatac ageacaceaa tggatettga etetttatte aattigeeag tetgtgtett
ttaacggggg catttagcct gtttacattt aaggttaata ttgttaigtg tgagtttgat
                                                                    1800
```

```
cctgtcatta tgatgctagc tggttatttt gcccgttagt tgatgcagat tcttcataat
                                                                    1860
                                                                    1920
gtcaatggcc tttacaattt ggtatgtttt tgcagtggct ggtactgctt ttttcctttt
                                                                    1980
tglattlagt gctlccttca gaagatcttg taaggcagga ctggtggtga caaaatcttt
                                                                    2040
cagcattige itticigiga aggattitat ticteettea ettatgaage itagtitgge
tggctctgaa attctgggtt gaaaattctt ttctttaaga atgttgtgcc aggcaccgtg
                                                                    2100
                                                                    2160
gctcatgtgt gtaatcccag cactttggga ggctgaggct ggcagatcac ctgaggtcag
gagitcaaga ccagcetgac caacatggga aaactecate tetactaaaa atacaaaatt
                                                                    2220
                                                                    2280
agccagctgt ggtggcacat gcctgtaatc ccaactactt gggaggctga ggcaggagaa
                                                                    2340
tegettgaac ecaggaggte aggttgeggt gageegagat ettgecatea taeteeagee
tgggcaacaa gagtgaaact ccatctcaca caaaaaaaaa aatgttgaat attggcccgc
                                                                    2400
                                                                    2460
activities ggottglagt gtttccgcag agaaatccac tgttagtctg atgggettec
ctitgtggat aaccegacct ttetetetgg etgecettaa egttittite atteetitea
                                                                    2520
accttggtga atctgatgat tacgtgcctt ggggctgctc ttctcgagaa gtatctttgt
                                                                    2580
ggiggicici giciliccig aactigaatg tiggicigic tigciaggit ggggaagitc
                                                                    2640
                                                                    2700
tectggataa taleelgaag agtgttttee aacttggtte eatteteece ateattttea
ggtacaccag tcaaacatag gtttggtctt ctcacatagt cccatatttc ttggaggctt
                                                                    2760
tgltcattcc ttttcattca tttttctcta atcttgtctt catgctttat ttcattaagt
                                                                    2820
tgatcttcaa teletgatat cettttttee aettgatega titggetatt gatacttgtg
                                                                    2880
talgetteae aaagttetig igetgigtti iteageteea teaggteati gatgattite
                                                                    2940
telagacigg tiaticiagi tagcaattet telaacette titeaaggit ettagtitee
                                                                    3000
                                                                    3060
ttgcagtggg ttagaatgtg ctcctttagc tcggaggagt tacccacctt ccgaagccta
ctlctgtcaa ttcgtcaaac tcattttcca tccagttttg tttccttgct ggcgaggagt
                                                                    3120
                                                                    3180
talgalccct tggaggagaa gaggtgttct ggtttttgga attitcagcc ttcttgtgct
ggillitect catelecting gattlateth cetting tt ttgatgttgg tgacetttgg
                                                                    3240
                                                                    3300
atgggglili igigiggaca tegilittgi igatgtigat getatteett teigttitt
                                                                    3360
agiliticie etaacaggea ggetietete etgeaggeet geiggagtit geiggaggte
                                                                    3420
cactecagae cetgitigee igagiateae tageagaeae igeagaaeag caaagaitge
                                                                    3480
tgcctgctcc ttcctctgga agtttcgtcc cagaggggca cccgccagat gctagtggag
ctctcctgta tgaggigict gtiggcccci gcigggaggi gictcccagi caggaggcac
                                                                    3540
aggggtcagg gacccactig aggaggcagt cigiccetta gcagagitig agigcigigc
                                                                    3600
                                                                    3660
lgggagatic geigetetet teagageigg caggeaggaa cattiacgie igeigaagei
geacceacag eegectette egecaggice telgiceeag agaggiggga gittiateig
                                                                    3720
ttageceetg actggggetg etgeetttet tteagagatg ecetgteeag agaggaggaa
                                                                    3780
tetagagagg eagtetgget atggeagett tgeagagetg tggtgggete tgeecaatte
                                                                    3840
gaacticeca gaageiitgi tialacigig aggggaaaac cacciacica agccicagia
                                                                    3900
atggtggacg cilcicecca caccaageti gagagiceca ggicgactic agacigetgt
                                                                    3960
```

gctggcagca	agaatttcaa	gccagtggat	tttagcttgc	tgggctctgt	ggcggtggga	4020
tccactgatc	cacttggctc	cctggcttca	gttccctttc	caggagagtg	aacagttctg	4080
tcgctggcct	tccaggtgtc	actggggtat	gg			4112

<211> 3408

<212> DNA

<213> Homo sapiens

<400> 2015

tteatectae tittgateea eteattaata acaettgget eageaggtee agggeacaaa 60 aacggtttca acaagtagca cgcaaggtca tgattcaggg acgattattc aatatgctga 120 gtgctgttcg tgaaatggac aaagagagta tactgagaaa gattggccaa gcaaaacaat 180 cgatagcaca agaggcgaat ttetteaaat tetteetgag geggateagt caggatgatt 240 300 ataccageeg giteteigtg tegeceaagg aggigeigee ettegetite eeagacigea geceaeceea ggaeteeaac gagttggete etgatggeet tggaetggte eeaattaagt 360 ctleagaagt teaaateaag cagagttatt cettetteaa tetgeaggtt ceteaactgt 420 acaaaattaa gagatatcag ccattctctg tccacaagtc ttcaacaagt tacagacctc 480 540 aaaagettge eegageeeta aageaaggag etgaggatga agteaceace ateacageee tteegaaaca ggaeteeaca acteagetet etggeaaaac ateaatettg ageatgaaac 600 cacctgagge citagecatg iciciagatt atgatectet gtatgittit aateccaace 660 caggattatt tgclgtaalg calcetelga cetalgeaga aacgltgata gattaceate 720 talgetetea ecceaagiae aaatteacea aagagteeeg eeaegggiee ageatteetg 780 tcacccaaaa gcagttictc catcacacgg acattattcc cggaataatg cactggaaaa 840 gettecagte cetggttete tectecetge eggacecete caagatggag accacaaaga 900 gelgegalte elicabilea illatgette egalagaegt ceelgecale eligatgeet 960 taccagaaga ggacagacta gaaacagtag aacgtgagct ctgtgagcag aatgtagaag 1020 ttatgitgac tecagaaatg atcaaagtgg aatteeetat gttgaactae aaggacatea 1080 ggaaggagaa agaagtgaaa gatcaagcac aaccagcaga gaaggccgga gagaagctgc 1140 tcgaggagat gaggaacctg cggggcaaag cactcaacac atacctgatt ctagaatgaa 1200 agtcaccagt aggttgaaaa ggtcgtggcc ccttggaaag attgtattga ctgtgttggg 1260 gatelggtge cacciggtgg atgecacaag aaaggeetet cetgacteec aagttgtaac 1320 cegiticeae canategaci iceanatani attiateaga teateateig igetitieti 1380 cettgtttca gaccactttt agglggaaaa ggcaaagaag gettatatgt attttettce 1440 ataatgagtc catcagaaaa agttccttcg gtgaaatcgt tgaccacgtg atgtttgggg 1500

actecetatg ggatcaa	lca tccgggttcc	ttagagacca	tggccataat	caggggctgg	1560
ccaagggaat gagtatc	ect gggttcaaca	gctgtttctg	aagacctgcc	agttcccctg	1620
tcttgcatta actcggg	tta tcatgccatt	ctccttctaa	ggccaaagat	acctgtaacc	1680
aaagaatcag gatactt	cac tgcagtcact	tcatttttt	ttcttttggg	gcagggtctt	1740
gctctgtcgc ctaggct	gga gtgcggtggc	acggtctcgg	ctcgctgcag	cctctgcctc	1800
ccgggttcca gcggttc	tee tgeeteggee	tctcaggtag	ctgggattac	agggacccgc	1860
caccaegece ggetaat	ltt tctgttttta	gtacagatgg	ggtttcacca	tgttggccag	1920
gcttgtctcg gactcct	gac ttcaggtgat	ccaccggcct	cattcccaat	ccatctccat	1980
teegeeatet tgetgee	cca tgggtaccca	cccttcccac	tgtgggcaac	catctcttta	2040
gtttctggtt tatcctt	ett gtgggtaatt	tttaaggcct	ctcggggtgc	tgggattgcg	2100
ggcgtgagcc accatgc	ctg gccaagcagc	ttcattttag	aagtgattat	tattgctttc	2160
ctttctagaa cttcagg	ttt gtgaagtatt	ttctcaatga	tcctcaaaac	attctaagac	2220
ataaagtagc tgttatta	agi gigalittai	gcagaaactc	aggcccagaa	agcttcatgg	2280
acttacccaa ttagcaga	agg agccaggttt	gggcaggatc	ttggtttcct	gcaaaggttt	2340
cgttgcctag ccaggcg	tgg tggtgtgtac	ctgtagtccc	agctacctgg	ggggctgggg	2400
tgggaggctc acctgage	ccc aggtagtcaa	ggctgcagtg	agccatgatc	ctggtaccca	2460
gtccactctt ctctcta	cta catggtaatc	aatgaaaata	ttacagattt	acattttta	2520
actttttatt taaactt	tca gctttggagt	ctctaagagt	aaagatatta	tgtgatgata	2580
tttgtatttt acttaat	tgc ttattcttta	aaacatgtaa	tatagaaaaa	aatacaaatt	2640
agcaaatgte ctttgct	cta aagaaatcag	ctggcaagtt	tgccccaccc	agcagcagcc	2700
atgtcttgct catttct	gta tecceageat	gcagcaagat	gtttggcaca	atgcaggctc	2760
tcaataaatg ttttttg	agg cigggiatgg	tggctcacgc	ctgtggtccc	tgcactttgg	2820
gaggctgagg caggtgg	atc ccttgagccc	aggagttcgg	ggccaccctg	ggcaacgtgg	2880
tgaagacctg cctctac	aga gagcacaaaa	gttggccggg	cgtggtggcc	catgcccagc	2940
tacttgggag gctgagg	tgg agggatcgct	tgggcctggg	gggtcgaggc	tgcagtgggc	3000
cgacattgtg ccaccgc	act ccagcciggg	cggcggagca	agaccctgtc	tcaattttt	3060
aaaaattggc taggtgc	agt ggctcatgtc	tgtagtccca	gcaccttggg	agaccgaggt	3120
ggacagattg cttgage	tca ggcattcaag	accageetgg	gcaacatggc	aaaaccccat	3180
ctctacaaaa aatacaa	aaa agattagcca	ggtgtgttgg	tgcacatctg	tggtcccagc	3240
tactggggag ggtaaga	tgg aaggateget	tgaccccagg	aggctgaggc	tgcagtgagc	3300
caagattgtg ccactgc	act ccagcctggg	caacagagca	agaccctgtc	tcaaaacaat	3360
agcaataatg titging	aat taaggaatat	aaaagaaatg	tgaaaact		3408

<213> Homo sapiens

(400) 2016						
gaagggctgc	tggagcgcgg	ccagaacgga	cgccgaggcc	gaggaggcgc	cgagagcgag	60
tgagagctgc	tagccagttg	tcacctctca	cagagaggtc	cacatttgct	gaaatgtaac	120
tcttatctca	tgcactggga	gtgatgaact	tcactaggaa	atcatcgttc	tttctggaat	180
tagacgatat	aagctgcagc	tcagaatcag	agcaggtcaa	gttgcttctt	ttcagtgatc	240
ataatcaatt	atccaggaaa	agggacagaa	gaaatcagga	aaaaggagaa	tagactcttt	300
atgcatagga	gctttaatat	atagttgaca	cttgaacaac	atggatctga	actacacaag	360
tccacttatt	ggtggatttt	cttcagcctc	tgccaccctt	gagacaacaa	gaccaatacc	420
tcctctttt	ccttctcctc	agcctactca	acttgaagat	ggtgagaatg	aagaccttta	480
tgatcatcca	cttctacttg	ttgaatagta	aatatattt	ttccttatag	ttttcttaat	540
aacactttct	tttctcttga	ttactttatt	gccagaatac	agtacatagt	acatataata	600
agcaaagtat	gtattagttt	actgittaag	tgataggtaa	ggcttccact	caacagcagg	660
caacagccag	gtgttgtgac	acatgcctgt	aatcctagca	ttttgggagg	ccgaggtgga	720
ggatcgcttg	agcccaggag	ttcaagacca	gcctaagtaa	catagtgaga	ccccatctc	780
cacaaaaaaat	taaaatacct	aatcatggtg	gtgcatgcct	gcaatcccag	ctactcagga	840
acctacaata	ggagccaaaa	aggtggaggt	tacaatgagc	cattattgca	ctactgcact	900
gcactcctgc	ctgggagaca	gagtgagacc	ttgtctcaaa	acactacaac	aaacaacaac	960
aacagcaaca	aaaatcagta	ggtattaata	gttaggtttt	ttgggagtca	gaagttatac	1020
acagattttg	actgtgcagg	ggatcagcgc	tectaaegee	tgcattcttc	aagggttacc	1080
tgtaticttg	atacaaattc	tectteagat	ttaagtatta	tagatattit	tccagtctat	1140
agcttaccta	ttcattttct	taataatgtc	ttttgattga	tttttaattt	ttaactttgg	1200
tgaattccag	ttgtatactt	ttttttatga	ttagcatiti	tgtgtcctat	gaaactgttg	1260
ccttcctcaa	tgtcactaaa	ttctcttagg	ttttcttcta	gcaagtttat	gtttcaaatt	1320
ticaccetta	ggtctataat	tcatcccaaa	tttatttttg	tctgtaaagc	aatgtcacaa	1380
ttcattttt	ticicaatat	agttacccag	ttgtttcaaa	actggitati	aaagttttic	1440
tcttaatcat	tgaattttct	tggcaccaaa	ttattaactc	1tgacaaaaa	taattgaccc	1500
ttaagtaagc	agacagacaa	gcagtgcttc	tattttatag	caatgtaaat	aatacacaac	1560
ttacacaaag	actttttaga	agctaactaa	cagtggtcct	atctaagtac	gtacaccaga	1620
ttttttataa	ccactttaa	aataaaagta	tttagattti	aacacataga	ttaggacaga	1680
gaaagcatat	ggtggaataa	actgtatctt	tttggccaga	tggtgctatt	tctaggtcat	1740
cttgataaag	agaggaggca	aacatgaaaa	cttaatgaaa	aactatttat	gatgctggag	1800
agaacatctt	ggctttgagt	cacttttaaa	tcatagaaga	ggattattcc	ataaaattat	1860

ttataatgcc	taaaattatt	ctttgcccaa	atcataaatt	ttcaggatta	ccaagaaacc	1920
atttagtatg	tatagagtgt	ttcagcaagt	gcagagatgc	ccaggtggtt	gggattcaat	1980
acatcgagct	gtcacgctgc	acattcttgg	agtacaacct	taatgggcat	tttcccacct	2040
gtgcgattcc	tctgttttca	ccccactcca	ttcatatttc	acaaactact	ctaattatag	2100
tatttattat	tgacctcagg	aaaaagaagt	ttgaaagggt	ggaaaaaaaca	tgcattttgt	2160
ctccatggat	agtaaatcac	tgagctattg	ttccttggga	atcccaattc	atgagaaatt	2220
acatagactt	ttgccctaac	actaatcagc	tgcctgatct	gtaaatattt	cagctccttg	2280
cctgtatcta	tttctccttg	cagaaaactg	taatttatct	agatttttct	aataattcac	2340
tgacatttta	ctgctagcca	atgagtaaat	cattgttgct	tttggtatct	tatgattttg	2400
ttcttttgtg	tcaaagttta	gctagtttca	tctatcaggt	tggaataaaa	aatgcaaatt	2460
atgactatac	cacttatata	gttacatgat	ctactgacca	aagttaatca	tcactttaat	2520
cttggtaact	cattcagage	cctaattgta	atagactttg	cctgagtcac	ctagagagtg	2580
gtctcaataa	tcccctttta	tttttcatgt	agagaaaagg	gcacacaaaa	tgatattatc	2640
tcgatcaccc	agcacatgta	ttaaactata	acagactttt	taaatcatgt	gtgatctttt	2700
attttttgac	tgaaagggac	taagtttgct	gcccagagaa	gtctttaggg	agcaaggaaa	2760
ggtaagcaaa	taaacttatc	tggagtcaaa	ggtctcaagg	aaaatcttgc	tttctataaa	2820
aggcagacaa	cgtcaagact	catagatttt	cccagggcta	aaaatcagag	ccaattgcct	2880
cccatcttga	aaagactcat	tcatcatgct	ggttgaagta	tcacagatct	tgtcaaaata	2940
ttcatgactc	acatacgacc	catccaaaag	acaaaagcca	acaaaatatt	ttaccaaatt	3000
ctaaaatagt	gtttgtttta	ttattctttg	ttattcttca	acaattattg	ctacctttac	3060
tatatgaaat	ataatagcaa	ttccttgtct	tcatggtctt	tctgttacag	acatgiitta	3120
cactgattat	accactttag	tgaaattcat	catacatatt	cctgatccaa	attccttttt	3180
tattaaccat	atatgagaga	aagtggatat	taaaataatt	tigaiggtaa	aataagcgaa	3240
aaaataaagc	aagcatggtt	aaaatgatta	aattgtggaa	aagtgaccca	tgtgtttcag	3300
ataaactgac	gcttgagggt	ttttgttgtt	attgttgttt	aaattttatt	ttattttaat	3360
tttaagttcc	aggatacaag	tgcaggatgt	gcaggtttct	tacataggta	aagatgtgcc	3420
gaaatggtgg	tttgctgcac	ctctcaacct	atcacctagg	tattaagccc	tacatgcgtt	3480
agctccctcc	cactgcccct	gcagcagatc	ccagtgtttg	ttgttccctc	cctgtgtcca	3540
tgtgttctca	ttgtccagct	cccacttgta	aataagaacc	tacggtgttt	ggttttctgt	3600
tcctgtttta	gtttgttgag	gataatgact	tccatgaagc	ttgagttttc	attctacaat	3660
ttactgaatg	acatttgagc	agctagctga	ctttttaatg	cctigatitt	aataattcaa	3720
tgagttattg	ggtgagataa	tttagaacag	catacatgat	atcgttatta	ttagtcaata	3780
aaatgctatt	tatcttattt	attactcata	acaaaaatat	gtatatgacc	cttcgctatg	3840
tttgaatatg	tgatatattg	aattgaattc	actgtgaggc	ttcagtaggt	acctataata	3900
ttcaaatatg	ttacctgaaa	gctgtgaaaa	atatatttt	aaaaattag		3949

```
<210> 2017
<211> 3618
```

<213≻ Homo sapiens

60	cggccgggga	tcgggcgctg	ggccgggaca	ggccagtcgc	ggatcccggg	gagagtccgg
120	ccagtagatt	gccagaggga	agctttctga	aatatggcag	agatagacag	cccgctgctg
180	caagcaagca	atttagagtt	aactcatgaa	ctggagtccc	atgtgtagct	ggaaggaacg
240	attgatgctg	gagacaagtt	aacagcttga	gagaagatgc	gcttttagca	agatacgaga
300	ttaaaagcag	ggaagataaa	tacaagttat	tttcaacagg	agaaaaaagct	aacgtcaagc
360	gatctggagt	taagtgtcaa	gattatataa	tcagagacaa	aaccagtgaa	ctaatattca
420	gaagagcaga	attgcaactt	aaaacttgga	gacgtcattc	ggaaaaaagat	cgctaataca
480	ataaaagaat	agcagctaag	tagaagagaa	gctaaaataa	aatacaagaa	aacaaataag
540	ttgatcaacc	gaatcttcgt	tggagaatca	gagctggaat	taagttaaat	gggtaacagt
600	ggaaagaagt	acaagttcaa	agtcaaaact	agaacaatgc	tgaagagata	aaaaccaaac
660	ttgacctttg	cctgagcagt	aaggccagcg	aagctttcgg	ctctacacta	catccactgt
720	gaaatgagca	aaaatctgag	ctcaagtagt	aggagtcctc	atctcgagca	ggtgcttttt
780	atggaaattc	catggaagaa	aaggaaaaga	gagttcactg	gaaagaacct	agatatcatc
840	acattgcatc	aggccagaga	aaaacaacag	caagttctag	tgttgataac	cagaaaaagtc
900	acagatggtg	aagctttgcc	aaacaagaac	cagaatcgga	tggctcagaa	aaaccccttg
960	gaagacggga	ctctgatgag	gtgactggag	gctccagtga	gaattctggg	gcatctccca
1020	tctgaggaag	cagtcacaca	ccaccctctc	agatgcacat	atccaagtcc	.gcagaggaag
1080	gacagcaget	agcatctgat	tgtatctgac	ggaagtgaaa	tagcaggatg	gggtccagtg
1140	ctatattctt	acacaagaag	agagaccaga	tttggcataa	ggaagagact	ctatatttga
1200	aattctgaat	tggaaaggga	atagtcctct	aaagctctaa	ggcacagtgg	ggcagcagga
1260	caatcccaga	taaaaaattt	atgaactgaa	agttcctcgg	ggaacaagat	taagtaaaaa
1320	ttgaccccag	aagccctatt	ссаасасссс	tcgagtgaag	ttcatcttca	gactcgatta
1380	glgccltcat	aacacaatta	ctggaaaagg	aactcactct	aaagcatcct	ctttaatgcc
1440	gtagcactag	cagtataagt	ctaatgtttt	ttaaggattc	acccccaaag	cacacctgcc
1500	acaaatagaa	gatgtttggt	gctctgacag	ccacagttaa	cttaagccag	ccaaaaggca
1560	ctagttgatg	tgatcttgat	ctcaggaaac	ccactgagac	catgatacga	acgctataag
1620	ttattttcct	tgatgatgga	acacgagttg	gagaatatgg	agaagtttta	gagacagtac
1680	gcaaagaagg	ctgtgaccca	accaggaaca	aattcagatg	ggactctcca	atgactcctt
1740	gaaagcagaa	tccttcttgg	tgcaccgttt	actectecee	caaacctcca	tggcatacag

tttatgctgt	agccaaatca	ggtattcgaa	tgtctgaggc	cttcaacatg	gagagtgtta	1800
ataaaaattc	tgctgcaacc	ctttcctata	ctacatcagg	actttataca	tctctgatat	1860
acaagaacat	gaccacccca	gtgtatacaa	ctttgagggg	aaggcgaccc	aaataagtag	1920
cagccctttc	ctggatgact	catctgggtc	agaggaagaa	gacageteca	gatccagctc	1980
ccggacgtca	gagtcagact	cacgcagtag	gagtgggcca	ggcagcccca	gagccatgaa	2040
acgaggtgtg	tctctctct	ctgtggcttc	tgaaagtgat	tatgctattc	ctcctgatgc	2100
ttactccaca	gacacggagt	actcacagcc	agagcagaag	ctcccaaaaa	cttgctcatc	2160
ttccagtgat	aatgggaaaa	atgaaccact	ggaaaaaatct	ggttatttat	taaaaatgag	2220
tggtaaagtc	aagtcttgga	agcggcggtg	gtttgttctt	aaaggtggtg	aattacttta	2280
ctacaaatct	ccgagtgatg	taattagaaa	accccagggc	catattgaac	ttagtgcatc	2340
ctgtagtatt	ttaagaggag	ataacaaaca	aacagttcag	ttgaccactg	aaaaacacac	2400
atactatctg	actgcagatt	ctcccaatat	attggaagag	tggattaaag	tgttacagaa	2460
tgttcttcga	gtacaagctg	ccaacccact	ttccctgcag	cctgagggca	aacccaccat	2520
gaagggattg	ctcactaagg	taaaacatgg	atattccaag	agagtctggt	gtacactaat	2580
aggaaagaca	ttatattatt	ttcggagtca	agaagataag	tttcctttag	gtcagatcaa	2640
actctgggag	gctaaagtgg	aagaggttga	cagatcttgt	gattcagatg	aagattatga	2700
agccagtgga	cgaagtctgt	tatccacaca	ttatactatc	gttatccatc	ccaaagacca	2760
aggtccaact	tacctcctaa	ttggatccaa	gcatgaaaag	gacacttggc	tttatcatct	2820
gactgttgca	gctggaagca	acaatgtaaa	cgttggatct	gaatttgaac	aactggtttg	2880
caaattgcta	aatatagacg	gggagccttc	ctctcagata	tggagacacc	ccactttgtg	2940
tcacagtaaa	gaaggaatca	tttcccctct	gacaactcta	ccttccgaag	ccctgcagac	3000
agaagctatt	aaattattta	agacctgcca	gctttttata	aatgctgcag	ttgactctcc	3060
tgcaattgat	taccacatat	ctttagccca	gagtgctttg	caaatcagcc	tgacacatcc	3120
tgagctgcag	aatgaaattt	gctgtcagct	tattaaacag	acaagacgaa	gacagccaca	3180
gaatcaacca	ggaccattgc	agggctggca	gctcttggca	ctctgcgttg	ggctcttcct	3240
tccccatcat	cctttcctgt	ggctcctcag	gcttcaccta	aagaggaatg	cagattccag	3300
gtgtgcagaa	tactagccag	ctgaactgtt	tatgtggcct	ctgaaagtct	acgataaatc	3360
ataagtattt	aacgatctgc	caggtacatt	ttcagaagaa	tgtatgaaac	aaatattggt	3420
acaggaagcc	tttggttatc	attgatgtgg	agctaggaaa	atatttcctt	tgttatgtta	3480
aatctcttag	ggaagattgc	aataaatact	tgaaaaactg	acagagaata	tttttaagtg	3540
aaaagtgcat	ttgcatttca	agtatgaatg	acttagcatt	agtgggtgtt	cattcaataa	3600
aagcaactat	tttgtttc					3618

<213> Homo sapiens

<400> 2018

agttgaagtg ttcactgata agtatgttaa ctaatgatcg agacagtaac gaaaaatgct 60 ggcactggga tictctccct tcccagacct acctgctggt atticctggg accttgaccc 120 180 tgccccaccc cctcagccgt gcccatctct gcagactccc agatcacatc tgggctgatg ggctggccca ggcctgtcta tttttcagtt cccaattaga agtctagaac ctgacaaclc 240 caggagttct tgggaggacc agtacaacgt tctaaaaagc ctgagacgcc ttacaaaaag 300 caagtateat ttggagtaea atteetaate tgtteatgte etgetgaagg agggaaggag 360 ggagaggaag gcaggggagt tgatgcattc atataacaaa cactgctggg tgtctgggtg 420 480 cccagagcaa agctgggcca ggccttcacc agatcaagcc ccacagacca gctggtgccc atgcgctgct ggtggtttgg ggcctcctgt tcctcctcta gctgggagta atcacagttg 540 600 telgacelga ticcaacita aggiceceae teletigece calcaagaat eeetgattat ttacttttcc ctagaaaatc tggggaaatt cccacatttt aattttgcag cagaatcttt 660 tgagcagett ttggaaceae agtgtttgee aagataagag tttgagaate cagcageeet 720 780 gggtgcctgg ctgaatttgg tttcctgcat gtgctgggtg tgggcggggc cacgcacagg 840 ccctgcatgg gaggactcct caccccaggc ctgtggtgct gcagacaacc gtctcctgtc 900 tacactgcga cccagccaca agctgtgggg tctcagtggc ctggggggaa gcagctccac 960 teteetgeee tteetggetg eccetttggg tteeageegg ggteaegtee ageeteeaet gggaaaccag tgactgaggc ctggacccag aggtggacca ggcatctcct ggccacctgt 1020 gacctgggaa gaagcgagtc agtggcccgt tcaacctgct ctgcagctgc tataaatagc 1080 1140 ctccctgttt ccaagaggag gtaaggaag! gtltatcttc taaaaaccag acgtttcctg 1200 atgetetgag egitaeteag tgetaeagag gagatgeaea egiteeeeact atgittetgie 1260 ttgagaaggg gacaagagaa agaggaaaag gagccactgt actttatttt gcacctacag 1320 cgtgccttgg cactgggcta gagaggcacc ttcctgcgtg aatcctgtgc ggcaggtctt attgccataa taagtcacat caaagacact gctggtcata aaacactgtt ttacatacca 1380 tagggaaaaa cgctgccaat ettaactaag atgctacaac tgtacagtte ettecaatea 1440 gagatgttca cgtgtgaaaa aaaaactgtg ctacttacaa tctatgaaag ctggtgttat 1500 cccacttgge aggtaaggaa actgaggtee tgtgagtgaa gtgacctcat gatcacacaa 1560 1620 caggagatgg cagggctggg attcaaaccc gggagtgtct gctgccacat cccacactcc cactgcctgg ctccaagtcc caggaagctc gagactgtga gttttctccc ttgaaactca 1680 1740 cctggagaga gtccgggcac ctgtgcctat gtggagggtt ccagccccag ccaggcccct ccgctgccca caccctggga ggagaagcgg cctcccttcc aggctcatct gctcactgcc 1800 1860 cgcattctcc tggcagagct gaggtctgag agatctggac tccaacccaa gggccctctc 1920 tigitatica ggggtgicca cagitaggaa gggacciggg gccitgiccc accaccitcc

```
1980
taggccccgt gatcaccacc ccctcaagcg gggccccagc cccctgagcg ccccctcacg
                                                                    2040
tgacccagcc ctcggctgtt ccaggctcac tgcccatggt gtgctcttct gggccacagc
                                                                   2100
agccaggget ccagggcgag gacaggggac acctgaaaac acccegttgt tcatggtctt
gtgcccattc attcggagac tcctgaaaaa ctgggctgtt tgcaaagcaa atccagctcc
                                                                   2160
ttgtcctagc aggttctcag aacggggagt cccctgggat ggagctgctc ccctcacggc
                                                                    2220
                                                                    2280
agcaccacgt ttccagtccc tcgatgccac taatcagcat ggactgtgtt caggacacag
                                                                    2340
ggtgaacttt tetetgaece eeggtgetgg teetgtgeea geaegtagta gttaeteagt
agaggtttgc tgagtaagcc agaaatcaga ttatgagtgt tcaggggttt gataaaacag
                                                                    2400
caccacataa cgcacacaaa gatactccag aaacatttgc tgagtaccta gtacgtgtga
                                                                    2460
ggtgctgtga ggatagagca gagaggactg tgccccagct gtgatgctgg cagaggtgac
                                                                    2520
                                                                    2580
actaagaggg aaatgagata tttggggcag aatccactgg gctctcttgg ccatccgctg
ccttgggtct gttgaggtgg gtgcccaaag gctgccttct tgaccagaac ctgctgtgcg
                                                                    2640
                                                                    2700
etteaeagaa eeteetette attggaaatg etgggeacat tgeagteagt gagetgetge
                                                                    2760
caaaacggcg ttaagtagaa ccccagagg ccccgccggt tggtgatcac cctcaggtcc
                                                                    2820
tgccagggag acacagtgag gaggttggct aattgctgct ttcaggccct ggaaatcagt
cgccaaggcc caggagaacc ccggtgagtc cgtccagttg aggcagaggc aataacctcc
                                                                    2880
                                                                    2940
cattgctcgg ccctgcgcct gccccagtcc tggcaggggg caccggctca ggaacatgcg
gcctcctggc atttctcggt atttaactgt ctcgctgtct tatccgagtc cctaatgaaa
                                                                    3000
                                                                    3060
cgacttgtgt gacaatctgt ctgtgcctta cgaaagtgtc tgtgcacttt ttatcctttt
                                                                    3120
taaaagcaac ttttaaaagt ggatggggag gggggctagc atgcgtggta gggttctaga
                                                                    3180
aatctgtggt catcgctgaa atcctttttg catcatgttt tttgatgttg gagtgatgaa
                                                                    3240
gigtacatec eccaceccae acaceactae etgigtacag acettitaaa acatgiette
                                                                    3300
tittictgat teaatactgi gaccieteeg ataeagieta ateetigggg aleigtaate
                                                                    3360
aaggttttaa aacctgggaa gtgggttggg aagggtttgc actggtcttg agtgttgtgc
                                                                    3420
ttttctgtgt tgtgtgtttt gatttttgtc tttttatctg ttttatattg acataatttt
                                                                    3451
cctgtttaaa aaaatacaac tttggcttgt t
```

<211> 4497

<212> DNA

<213> Homo sapiens

<400> 2019

agagctgggc cctgtgaccg cagaccagag ggagaggagg aggctggact gggctgcgag 60 tgtgggagag ggtggactca gggccccagc aggttagtgg gagatggaac aggcaccag 120

ggctgccaag	aaccccagca	aagccgggct	cccaggtggg	tggacaggtc	ccagagccag	180
tgagggccgg	ctcctcccat	gagggtggct	gcacaccccc	tcctgccggg	gcaggcagtg	240
ctggtctgcg	cccgctcccc	agccccccac	cggctgtgcc	agctgggccg	cagatggacc	300
acatggggaa	cagctcccag	ggggccccct	ggctcttcct	cacctccgca	ctggcccgag	360
gcgtctcggg	gatcttcgtg	tggactgccc	tggtgctcac	ctgccaccag	atctatctgc	420
acctgcgctc	ctacaccgtg	cacaggagca	acgttacatc	atccgcctgc	tcctcatcgt	480
gcccatctac	gccttcgact	cctggctcag	cctcctc	ctcggagacc	accagtacta	540
cgtctacttc	gactctgtgc	gggactgcta	cgaagccttt	gtcatttaca	gcttcctgag	600
cctgtgtttc	cagtacctgg	gaggcgaggg	cgccatcatg	gctgagattc	gtggaaagcc	660
catcaagcca	ctctgcagtt	ctgcctggtg	aagcccgtca	tggccgtcac	caccatcatc	720
ctccaggcat	ttggcaaata	ccacgacggg	gacttcaatg	tccgcagcgg	ctacctctat	780
gtgaccctca	tctacaacgc	ctccgtcagc	ctcgccctct	acgccctgtt	cctcttctac	840
ttcaccacca	gggagctcct	gcggcccttc	cagcccgtcc	tcaagttcct	caccatcaaa	900
gccgtcatct	tcctgtcgtt	ctggcaaggg	ctgctgctgg	ccatcctgga	gcggtgcggg	960
gtcatcccgg	aggtggagac	cagcggcggg	aacaagctgg	gggctggcac	gctggccgcc	1020
ggctaccaga	acttcatcat	ctgcgtggag	atgctgttcg	cctccgtggc	cctgcgttat	1080
gccttcccct	gccaggtgta	cgcagagaag	aaggagaatt	caccagcccc	cccggcaccc	1140
atgcagagca	tctccagcgg	catcagggag	acagtgagcc	cccaggacat	cgtgcaggac	1200
gccatccaca	acttctcccc	cgcctaccag	cactacacgc	agcaggccac	gcacgaggcg	1260
cccaggcccg	gcacccaccc	cggcggcggc	ggctccggcg	ggagcaggaa	gagccggagc	1320
ctggagaagc	ggatgctgat	ccctcggag	gacctgtagg	ggggcctggg	ctgccagtgc	1380
tgtagggacc	caggctgccc	aggcctctgg	ggaagaacag	ggtccccca	cccaccaact	1440
cctgccaaag	gtggggcctc	tcctgagagc	ccacctgtga	ggccctcgga	gcccacttcc	1500
catcctccct	ccagccaggg	ggtcagggca	cctgatggcc	ctggcaggca	cccaggtggg	1560
cccgccaccg	caggagaggg	cacctgagcc	aatcggaaga	gcctggggac	cccctgggat	1620
cacccagcca	tcagccccag	gagccactgt	ggggcggaga	gtgagtgtgg	ctgcggggcc	1680
ttggctgcac	ggaccccatg	ggagctgcga	gtgggtcaga	ctccctggtt	caggagacag	1740
acagcggacg	gatcccaggc	tgggcagctg	gagggagggg	cgccggggcg	ctgggcagcc	1800
gggctctgac	acagtcagca	gctccgggcg	ccgcaggccg	gcggggtcca	cacaggctgg	1860
ccggggctgg	gcctccttgg	agcctgctac	ggccctcgtg	ggcacgtgga	gaagggccca	1920
cgtgtctcca	cacgccagcc	acaggggagc	cctggccagg	cgcccagcca	ggggagcgtg	1980
tgcctgggat	gggtcacaga	accagcgggc	acctgtgagg	ctggccagca	ccgtggggct	2040
gtgggaatcg	ctcttattta	tatttaaaca	ccttggattt	tctaccgggt	cttggcttct	2100
gttcccgcag	ggcatgagcc	tgaggagcag	gacgcggtgg	gggtcacagg	aggctgctgc	2160
tcagagtctg	catgcgggaa	aggggtccca	cctgtctggg	gtgggcagcc	tcgtggtcca	2220
gggcagtgca	gggcagagcc	tgggctgtgc	gatcacagcc	actgcctttc	tcctgggagc	2280

ctccacttcc	tccaaaacgg	gccttgtgcc	agccccaccc	gcggcgagcg	gacaaggcca	2340
cgagggcagg	gccctgagta	cctgggcggg	ggggacactc	ccagggggca	cagagggggc	2400
tcccacctgg	gcacctgcct	cctgcccttc	tcttcttcct	ccacgtgcca	ggtggggccc	2460
tgggtttgag	gagcctcgga	cgcgtgccct	gcccgcagga	agctggaggc	gtgcaagtgg	2520
cctcggaaat	cgcggccgca	agaacagtag	ccgcccaggg	actaaggggg	cttctgggag	2580
gacacacggc	tggcccaggg	cgaggggtgt	cactgcaggg	cgcccccag	gcccagggcc	2640
cgtcagggga	cagtacggtg	acccggcctg	caggtggcag	tcagttctgt	gtgtctgggg	2700
cccacagcac	aggttgggtg	ggggctgggg	caggggcagc	agaagtgggc	aaggcctggg	2760
gggctcaggc	actgggcgtg	gagagcagac	aggaagetee	agtgggcacc	accccgggac	2820
cgcggctccc	acccgtgctg	cccccaccc	atggccacgg	tcaccaggaa	cagcgggacc	2880
tggggtctcc	gagggactca	gcagggcggg	cacagaccag	tggagtccgg	gctagagagg	2940
gccagctccc	agcctcttgc	ttcctgggct	gaggacatgg	ggatccaagg	ccagtgggtc	3000
tgcagggccc	agcccggctg	cctgataaga	taggccgagc	tcctccctgc	acggctgcaa	3060
agacgcccac	ctgtcttatt	ggatccccac	aggaatagac	ccaccaggcg	gccccgtgt	3120
ctcactctgt	cagcaggtcc	ccagggacct	gctgccgagg	ggcagtttct	ggaggctggg	3180
ggcactggct	gggctctagg	cctgctctgc	ctttgccgtg	gagaaggcca	ccccgatagg	3240
ggtcaagttg	ctcaaatctg	cgtttggagg	gtatgtggcc	gagggctccc	tttctggaga	3300
cccagacacc	gcctgggctc	cgggcggcag	aggctgaggt	gtcaggggct	gagcccctat	3360
gtcagcaaca	cctcaggcct	gcactttagg	acaggggaga	agtcagtttc	cgccaaatgc	3420
ccctcagac	cagccgagga	ctgtgccagg	aaactgacat	gctcagcgct	caagccagct	3480
gggacagcga	ccgagcccag	agagacggag	caagttgcct	gaggtcacag	agcagggact	3540
tggacaccag	gcagccggct	ccacagaggc	cctctctcct	ccctgcctcc	tgaccctcag	3600
acgcctccgc	cccacgggtg	aggetgette	tgcttctttc	caacacgact	cgaaggaaag	3660
ccctgagggc	cgagcccgct	ctgcgtggac	ggaaggcagc	gtggggcggt	ccaggccggg	3720
gctcaacctg	cctcgagggg	gagcgtgggc	gcatgtgagc	gggagggacg	gagactagcg	3780
tggttccagt	gtcgtcatcg	ctgctaaaaa	aggggtttcc	cggtgacagg	ccccgacaga	3840
ggagcaggcc	atgaggcagg	caggagccac	gtatctgggc	ccagcgcacc	egceaagete	3900
tctagcctct	cctggcctca	gtatecttet	ctgggagatg	gtccagctga	aaatccccag	3960
catccacaag	aaagggtgga	agccctgggg	gccctggcct	ggcccaggtg	caggctgcat	4020
ggccgggcgg	ggcggtgtct	cctttcacag	cttccccgtc	tgtccgcagc	ctccaggagc	4080
cccacacagg	gctggggctc	tgtgccccca	actcacaccc	gtcggctccc	ccaggaggag	4140
caggctgggc	ccagagccgc	agggtgggct	gcagggaggt	ctgacttagc	tggggaaagt	4200
gccatccctg	ccattgctag	tgacaagctc	gggctgctgt	ggccccagca	cagattcaac	4260
actcactgcg	ctacgtgcca	gctgttgcac	actcacctcc	acacccaact	cacaggaagc	4320
${\tt aaggctgggg}$	aggagggaac	tggccccagg	ccacacagat	gctgcgagtt	gggattatga	4380

tegggtgeag tggeteacae etgtaattee ageaettggg gaggeeaagg egagtggatt 4440 gettgageee aggagtetga gaeeageetg ggeaacatgg tgaaaceeea tetetae 4497

<210> 2020 <211> 4590

<212> DNA

<213> Homo sapiens

<400> 2020

60 accacacca gctaattttt gtatttttag tagagatggg gtttcaccat gttggccagg 120 ctggtctcaa actcctggct tcaagtgacc cgcctgcctt ggcctcccaa agtgctggga 180 ttacaggcgt gagccaccac acccagcccc attgtctttt ttttaagaca ctggttctca 240 ctctgtcacc taggctggag tgcagtggtg tgatcaaggc ttactgcagc ctcaacctct 300 tgggctcaag cagtcctccc actttagcct cccatgttgc tgggaccaca ggtgcatgcc accaagecee actaattaaa acaaattttt ttttatagag aataggatgt agetatgttg 360 420 cccaggctgg tcttgaattc ctgggctcaa gtgatcctcc caccttggcc tcccaaagtg 480 ctgggattac aggtatgagc tactgcacct ggtctctgtc ttctttttt ttttaaggct 540 ctigttagaa igccgigaac agiigticic aaciattata igicalicca cgggatiggi 600 ttcctgctgg cattccatgg tctccggggt cctctgcagc accttcctgg ccttttgtca 660 tgtggatgct gcacagctga ctccacctgg tcttgttgat ggacagtttg tttcatgatt 720 tctcttatga ataaaacctt cacaagccat ccttctctat gagagtgttt gcttggcacg 780 cattectgag cactgeeect gageagaeeg ectatgatet etaagettgg gtteegtgtt 840 gccaaagege cttetggtgg acteagecea ggaggagece atgtgeecea egetggeeat 900 ggctgtggtc atgggctgac tgcatgtgtc tgactgggcc ttcgtctgag actgcagtga 960 tttcgctcct cctctcagat ccgcaaggat gctctccggg cgctcaactt tgcgtacacg 1020 gtgagcacac agcgatctac catctttccc ctggatggtg tggtgcgcat gctgctgttc 1080 agagactgtg aagaggccac cgacttcctc acctgccacg gcctcaccgt ttccgacggc tgtgtggagc tgaaccggtc tgcattcctg gaaccagagg gattatccaa gaccaggaag 1140 teggtgttta ttactaggaa getgaeggtg teagtegggg aaattgtgaa eggaggeea 1200 1260 ttgcccccg tccctcgtca tacccctgtg tgcagcttca actcccagaa caagtacatc ggggagagec tggccgcgga getgcccgtc agcaeccaga gacccggctc cgacacagtg 1320 1380 ggcggaggga gaggaggga gtgtggtgta gagccggatg cacccctgtc cagtctccca cagtetetae cageecetge geecteacea gtgeetetge etectgteet ggeaetgace 1440 1500 ccgtctgtgg cgcccagcct cttccagctg tctgtgcagc ctgaaccacc gcctccagag 1560 cccgtgccca tgtactctga cgaggacctg gcgcaggtgg tggacgagct catccaggag

gccctgcaga	gggactgtga	ggaagttggc	tctgcgggtg	ctgcctacgc	agctgccgcc	1620
ctgggtgttt	ctaatgctgc	tatggaggat	ttgttaacag	ctgcaaccac	gggcattttg	1680
aggcacattg	cagctgaaga	agtgtctaag	gaaagagagc	gaagggagca	ggagaggcag	1740
cgggctgaag	aggaaaggtt	gaaacaagag	agagagctgg	tgttaagtga	gctgagccag	1800
ggcctggccg	tggagctgat	ggaacgcgtg	atgatggagt	ttgtgaggga	aacctgctcc	1860
caggagttga	agaatgcagt	agagacagac	cagagggtcc	gtgtggcccg	ttgctgtgag	1920
gatgtctgtg	cccacttagt	ggacttgttt	ctcgtggagg	aaatcttcca	gactgcaaag	1980
gagaccctcc	aggagcttca	gtgcttctgc	aagtatctac	agcggtggag	ggaagctgtc	2040
acagecegea	agaaactgag	gcgccaaatg	cgggctttcc	ctgctgcgcc	ctgctgcgtg	2100
gacgtgagcg	accggctgag	ggcgctggcg	cccagcgcag	agtgccccat	tgctgaagag	2160
aacctggcca	ggggcctcct	ggacctgggc	catgcaggga	gattgggcat	ctcctgcacc	2220
aggttaaggc	ggctcagaaa	caagacagct	caccagatga	aggttcagca	cttctaccag	2280
cagctgctga	gtgatgtggc	atgggcgtct	ctggacctgc	catccctcgt	ggctgagcac	2340
ctccctggga	ggcaggagca	tgtgttttgg	aagctggtgc	tggtgttgcc	ggatgtagag	2400
gagcagtccc	cagagagttg	tggcagaatt	ctagcaaatt	ggttaaaagt	caagttcatg	2460
ggagatgaag	gctcagtgga	tgacacatcc	agcgatgctg	gtgggattca	gacgctttcg	2520
cttttcaact	cacttagcag	caaaggggat	cagatgattt	ctgttaacgt	gtgtataaag	2580
gtggcccatg	gcgccctcag	tgatggtgcc	attgatgctg	tggagacaca	gaaggacctc	2640
ctgggagcca	gtgggctcat	gctgctgctt	cccccaaaa	tgaagagtga	ggacatggca	2700
gaggaggacg	tgtactggct	gtcggccttg	ctgcagctca	agcagctcct	gcaggctaag	2760
cccttccagc	ctgcgcttcc	tctggtggtt	cttgtgccta	gcccaggagg	ggacgccgtt	2820
gagaaggaag	tagaagatgg	tttgtgaagg	aagtctcgtt	tatgaagcag	cattgtttaa	2880
taaatgggtg	gaggccctgg	gtctgaggat	ggtccagtag	tgttggggtc	aggaatcact	2940
gagacagcaa	ccctgtggt	gactgtccac	tgcaggactg	ggtggggtca	gcacagtgag	3000
atatgttagc	aggtgtgctg	acagcagaat	gcaagtgacc	ttcatctatg	tctgtcttaa	3060
aggtctgatg	ctacaggact	tggtttcagc	taagctgatt	tcagattaca	ctgttaccga	3120
gatccctgat	accattaatg	atctacaagg	ttcaactaag	gttttgcaag	cagtgcagtg	3180
gctggtttcc	cactgccccc	attcccttga	cctctgctgc	cagactetea	ttcagtacgt	3240
cgaagacggg	attggccatg	agtttagtgg	ccgctttttc	catgacagaa	gagagaggcg	3300
tctgggcggt	cttgcttctc	aggagcctgg	cgccatcatt	gagctgttta	acagtgtgct	3360
gcagttcctg	gcttctgtgg	tgtcctctga	acagctgtgt	gacctgtcct	ggcctgtcac	3420
tgagtttgct	gaggcagggg	gcagccggct	gcttcctcac	ctgcactgga	atgccccaga	3480
gcacctggcc	tggctgaagc	aggctgtgct	cgggttccag	cttccgcaga	tggaccttcc	3540
acccctgggg	gcccctggc	tccccgtgtg	ctccatggtt	gtccagtacg	cctcccagat	3600
ccccagctca	cgccagacac	agcctgtcct	ccagtcccag	gtggagaacc	tgctccacag	3660
agcctactgt	aggtggaaga	gcaagagtcc	ctccccagtc	catggggcag	gccctcggt	3720

```
3780
catggagatc ccatgggatg atcttatcgc cttgtgtatc aaccacaagc tgagagactg
                                                                    3840
gacgececce eggetteetg ttacateaga ggegetgagt gaagatggte agatatgtgt
gtalttiltt aaaaacgalt tgaaaaaata tgatgttcct ttgtcgtggg aacaagccag
                                                                    3900
                                                                    3960
gttgcagacg cagaaggagc tacagctgag agagggacgt ttggcaataa agccttttca
                                                                    4020
teettetgea aacaatttte eeataeeatt getteaeatg caeegtaact ggaagaggag
                                                                    4080
cacagagtgt gctcaagagg ggaggattcc cagcacagag gatctgatgc gaggagcttc
                                                                    4140
tgctgaggag ctcttggcgc agtgtttgtc gagcagtctg ctgctggaga aagaagagaa
                                                                    4200
caagaggttt\ gaagatcagc\ ttcagcaatg\ gttgtctgaa\ gactcaggag\ catttacgga
                                                                    4260
tttaacttcc cttcccctct atcttcctca gactctagtg tctctttctc acactattga
                                                                    4320
acctgtgatg aaaacatctg taactactag cccacagagt gacatgatga gggagcaact
gcagctgtca gaggcgacag gaacgtgtct aggcgaacga ctaaagcacc tggaaaggct
                                                                    4380
gateeggagt teaagggaag aggaagttge etetgagete catetetetg egetgetaga
                                                                    4440
                                                                    4500
catggtggac atttgagcag cctgacctgt ggggaggggg tctctcccga agagtttctg
titttactca aaataatgit atteteagat gettgatgea etgitggaaa tgtgattaat
                                                                    4560
                                                                    4590
ttaatcatgc agataaacca tttaaatgtc
```

<211> 4110

<212> DNA

<213> Homo sapiens

```
60
ataaggetac ctggctggga ccacagatgg agtctcgctc tatcacccag gctggagtgc
                                                                     120
aatggcgcga tctcggctca ccgcaacctc catctcccag gttaaagcga ttctcctgcc
                                                                     180
teagteteet gagtagetgt gattaeagge gtgegeeate aeaeceaget aatttttgta
ttttttagta gagatggggt ttcaccatgt tggcctaact cctgacctcg tgatccgccc
                                                                     240
                                                                     300
atcttggcct ccgaaagtac tgggattaca ggtgtgagcc actgcacccg gcccaaacat
                                                                     360
ttctttttct tttcttttga gacagagtct tgctctgttg cccgtggctg gagtgaaatg
                                                                     420
gigcgattat agitcacigc agcetcaaac teeiggeelt aagegateet eccateeigg
                                                                     480
cctcccaaag tgctgggatt ataggcatga gccgcagcaa ccactcctca catttcttga
gcatctgtga tgtatcaagc cagatgctgg gcactgaggt tgcagaaggc attgttcctg
                                                                     540
tcttctagga gccccaggct agcagggaag acggatgtgt atagagttaa ccacaatacc
                                                                     600
                                                                     660
aggecteaac ticccgictg taacacaggi ggaccatget agaitgteec agectgeect
                                                                     720
gtgcticatt agccggtcaa cagatccatc tcaaatacct cccatgggta ctcactgatt
                                                                     780
getttaacee aaaceatgge actettgaag acttteeete aggaagetea aggaetatge
```

atccttctgg	gtcagaactg	gacacacagc	caccagtgct	ggacaatggc	ggcggctcag	840
ggacacactg	gagccctggc	ccctgcagag	ctcccagcat	gggtgggaag	agagatgcaa	900
aatgaccaca	cggcgggtga	ggaggagctc	cctcggtgcg	gctgggatga	gccctagaca	960
ctctcaatca	ccccacgat	gaccccttcc	cagaggtccc	ctcagtcatc	tgccctgaac	1020
caagctcttc	ctgatcctag	accetecace	ctccctctat	cttccagggc	ttggtgacat	1080
tccaggcaga	aatttctgac	ccttttactt	tggtccctcc	ctccccagcc	cagtctctgg	1140
tcaaactgga	ttcctggctg	ttcccagaac	gagctgcctt	tecceaectt	gccacctctg	1200
cccttgttct	ctctgcctga	atgtcctcct	tcactagcct	cgctgccttg	cacatetete	1260
ctgagggctg	tcatcccaga	atgagctgca	tttgtccagc	ctggcccacc	atctaccaga	1320
acgtcctcct	tcagcctgtc	ccactgcctt	gcaaaacttt	tctgggggac	ctgttcacaa	1380
tgccttctgt	agcatactcc	aagaatccgg	cgcccctgg	agttgtgcca	cacagcaccc	1440
ctttgcagtc	aagctccctc	agcaccacca	cctccaccct	ggaagagttc	cccttccctt	1500
tgaaatctca	tgggactttg	cacccactct	ggctttattg	gaaggctttg	tatgtctcca	1560
cagggtaaac	acccatttac	tggggtgatg	atgtctccag	gatctagttc	atgtttgtcg	1620
ttggtgactg	gccccaccca	gttctgggca	agcaggctgg	atcccggcag	gaacagagcc	1680
caccagccta	aacttccatg	gaggtggaga	ggggacaggc	ttctgtctct	ttttggctga	1740
aggtgcatca	tgtccaaggc	ccctcttcta	gccaagcaga	gaagctgggt	gataaggatg	1800
ggtgagagtg	ggtgatgtac	cccggagtcc	tggcctcccg	gctcctcact	cccctacgcg	1860
taactttatc	cggccaatgc	cgcaaagact	gctggtgagg	ccagatgcat	gagtgatcat	1920
actcacaaca	gtcgtgaaac	tgccagtgat	gaaactggta	aggacaagaa	atgacaataa	1980
tcaaggtggg	gtttctcgtg	gacgtttcca	agacttcatt	ctcaaattct	ctccctcagg	2040
gtccccaccc	tgtcctccca	cctaagcctg	gaatgagggg	gcactggcct	gtggggaccc	2100
tggtcttcag	gctcccaaac	ctggctgggt	ctggttgccc	cctggcctta	acctgtgaac	2160
atccagctgt	ccctgggctg	tgattcagtg	tctgtctcct	gggtgacctc	agcatgggct	2220
ttgaggaagg	ggagagagta	gtttcttctg	agactggata	gtgactcagg	gacccagggc	2280
tggggcctca	aaagtgcctt	tgttggcctg	ggctcaggaa	tccagagaaa	ctggtcagga	2340
ggaggcccca	gtgacaaaaa	cccctccctc	tgccccgcc	cctctgccag	agccatataa	2400
ctgctcaacc	tgtccccgag	agagagtgcc	ctggcagctg	tcggctggaa	ggaactggtc	2460
tgctcacact	tgctggcttg	cgcatcagga	ctggctttat	ctcctgactc	acggtgcaaa	2520
ggtgcactct	gcgaacgtta	agtccgtccc	cagcgcttgg	aatcctacgg	ccccacage	2580
cggatcccct	cagccttcca	ggtcctcaac	tcccgcggac	gctgaacaat	ggcctccatg	2640
gggctacagg	taatgggcat	cgcgctggcc	gtcctgggct	ggctggccgt	catgctgtgc	2700
tgcgcgctgc	ccatgtggcg	cgtgacggcc	ttcatcggca	gcaacattgt	cacctcgcag	2760
accatctggg	agggcctatg	gatgaactgc	gtggtgcaga	gcaccggcca	gatgcagtgc	2820
aaggtgtacg	actcgctgct	ggcactgccg	caggacctgc	aggcggcccg	cgccctcgtc	2880
atcatcagca	tcatcgtggc	tgctctgggc	gtgctgctgt	ccgtggtggg	gggcaagtgt	2940

accaactgcc	tggaggatga	aagcgccaag	gccaagacca	tgatcgtggc	gggcgtggtg	3000
ttcctgttgg	ccggccttat	ggtgatagtg	ccggtgtcct	ggacggccca	caacatcatc	3060
caagacttct	acaatccgct	ggtggcctcc	gggcagaagc	gggagatggg	tgcctcgctc	3120
tacgtcggct	gggccgcctc	cggcctgctg	ctccttggcg	gggggctgct	ttgctgcaac	3180
tgtccacccc	gcacagacaa	gccttactcc	gccaagtatt	ctgctgcccg	ctctgctgct	3240
gccagcaact	acgtgtaagg	tgccacggct	ccactctgtt	cctctctgct	ttgttcttcc	3300
ctggactgag	ctcagcgcag	gctgtgaccc	caggagggcc	ctgccacggg	ccactggctg	3360
ctggggactg	gggactgggc	agagactgag	ccaggcagga	aggcagcagc	cttcagcctc	3420
tctggcccac	tcggacaact	tcccaaggcc	gcctcctgct	agcaagaaca	gagtccaccc	3480
tcctctggat	attggggagg	gacggaagtg	acagggtgtg	gtggtggagt	ggggagctgg	3540
cttctgctgg	ccaggatggc	ttaaccctga	ctttgggatc	tgcctgcatc	ggtgttggcc	3600
actgtcccca	tttacatttt	ccccactctg	tctgcctgca	tctcctctgt	tgcgggtagg	3660
ccttgatatc	acctctggga	ctgtgccttg	ctcaccgaaa	cccgcgccca	ggagtatggc	3720
tgaggccttg	cccacccacc	tgcctgggaa	gtgcagagtg	gatggacggg	tttagagggg	3780
aggggcgaag	gtgctgtaaa	caggtttggg	cagtggtggg	ggagggggcc	agagaggcgg	3840
ctcaggttgc	ccagctctgt	ggcctcagga	ctctctgcct	cacccgcttc	agcccagggc	3900
ccctggagac	tgatcccctc	tgagtcctct	gccccttcca	aggacactaa	tgagcctggg	3960
agggtggcag	ggaggagggg	acagcttcac	ccttggaagt	cctggggttt	ttcctcttcc	4020
ttctttgtgg	tttctgtttt	gtaatttaag	aagagctatt	catcactgta	attattatta	4080
ttttctacaa	taaatgggac	ctgtgcacag				4110

<211> 3937

<212> DNA

<213> Homo sapiens

<400≻ 2022

aatgctgaga	cagactccca	gaagatctga	gcgagtcgcg	tagctgagcc	cggcaggggc	60
tggggtggtg	ctgctgctat	gagctgcacc	atcgagaaga	tcctgacaga	cgccaagacg	120
ctgctggaga	ggctacggga	gcacgatgcg	gccgccgagt	cgctggtgga	tcagtcggcg	180
gcgctgcacc	ggcgggtagc	agctatgcgg	gaggcgggga	cagcgcttcc	ggaccaggtc	240
aggcagaggt	atcaagagga	tgcatccgat	atgaaggaca	tgtccaaata	caaacctcac	300
attctgctgt	cccaagagaa	cacacagatt	agagacttgc	aacaggaaaa	cagagagcta	360
tggatttcct	tggaggaaca	ccaggatgct	ttggaactta	tcatgagcaa	atatcggaaa	420
cagaigitac	agttaatggt	tgctaaaaaa	gcggtggatg	ctgaaccagt	cctgaaagct	480

caccagtctc	actctgcaga	aattgagagt	cagattgaca	gaatctgtga	aatgggagaa	540
gtgatgagga	aagcagttca	ggtggatgat	gaccagtttt	gtaagattca	ggaaaaatta	600
gcccaattag	agcttgaaaa	taaggaactt	cgagaattat	tgtccatcag	cagtgagtct	660
cttcaagcca	gaaaggaaaa	ctcaatggac	actgcttccc	aagccatcaa	ataactgaac	720
tctgaatgat	ggctggagat	tgtctatcaa	ggaaggaagt	tactgtcttc	ccattcaagt	780
actgtccatt	aagtgtcttg	cctcagattt	gatttaatct	taattaaagg	tatcaggtgg	840
caatttagaa	ttccagtcaa	tattggctgt	ccacagttct	cagatgtgtt	aatgtgaata	900
ctacatgctg	aatttcacca	ttcctttctc	aaagagacta	cttttaattt	tcatttctgg	960
gaccttgatt	tatataaact	atgttttcag	ttctttgtta	tttttcacat	ctctgaaact	1020
ttgagcattt	tttataagcc	agcaatttat	tttacatagc	attgtaaaat	acacttctag	1080
gaaattttag	gaaagattta	actgtttaaa	tctatttggc	ataaaccttg	attttttt	1140
tccatttgac	aaaaataata	caattccaca	gaactagatc	agcagattct	ctgatttgta	1200
atgtcattca	cctgtgacat	tttaagtctc	tctggtgcta	agaattggca	ctttatagcc	1260
tggtgccttt	acttttaatt	tgagagaacc	tactgctagt	cccaggaaac	acacttggaa	1320
ataagtcagc	tattitttt	gcccagtgat	gctatagttg	tcatattgtc	caaagttcat	1380
attgttcaaa	gctgaggagc	ttgtcctgtg	tatgtgaatg	cacacatgtg	cacttagttc	1440
aaatactaaa	agtagctttt	attaaatata	atcagccaaa	aacacacaca	aaataaaaaa	1500
aacaaatata	agtagtcagt	ttttcaatgt	tatcctacta	gttctacatt	ctattttaat	1560
ttttatacaa	tttccatttt	atagttaaga	accatcactt	acttggattg	gatgtctttc	1620
attcctagca	ctaatagttg	gctttctttt	tttttgttta	catagaagca	gggtttttt	1680
ttatcttttt	tcttttttt	tgtttaagct	atataaaaag	gtgaggaagc	agttttgtta	1740
cctaatgaaa	attattacac	tcataatgct	gtgtaggcaa	cattgagatt	caaatgccca	1800
gtggtcaact	gggttcactc	atcaactcat	tecegtecea	gtttactcac	atttcaaatt	1860
tataaatttc	ticaigitat	actattctat	ttagatttgc	ccagaattag	ttgaaataat	1920
gctaaacctg	tcaatatttt	ccagtaacat	taagcaccat	actgcatggg	agagacacag	1980
tactaaaaaag	agttgttagt	gctttatgtg	agtgatattt	ctttcgtaat	gctataaaga	2040
actacagtta	aaataacaga	atatttaaa	gatgtcctaa	aagcatctga	tcccagtaat	2100
aactaatgga	tgtcatctag	agcagtgggt	gttaatgaat	aggtatatgt	catttaagaa	2160
tttttcaaat	ttctgtttga	tatcctgcat	agaatttgac	aaaaaaaaca	cttccaagtg	2220
tgagcatttt	ttatttcatt	tcccaagagt	aagtaagtaa	ctattagccc	agccatctgc	2280
ctcgaagtat	accttaagtg	accccataaa	tccattcaag	aggcaggtac	tctataccat	2340
ttggcagcca	cggccaaacc	taccatggcc	agatttcagt	gaaaatgatg	aagtaatcaa	2400
atcaaggtat	aatatggtgt	ccctttatgt	gctttatgtt	cctttagagc	tgtttataaa	2460
gttctttata	tctcaagtgt	taggataaat	cgacatacta	actiticccc	ctgcaaaatt	2520
aaaagcctga	ggtacaagtc	taagaagctt	ttagtgctct	acataatata	aattctggct	2580
ggtgttaatg	ctatgaagat	aatatgtagt	tagaaaattg	agtcggggag	gaatgctctt	2640

ctttttaagt	ggattttaaa	gtttctcctt	gagtggatga	agaacttgcc	tggtttgcaa	2700
aaatcttagt	tcaaaattat	attttctaac	aaaaactgca	ttttgagaag	ataagctaat	2760
tttactcagt	agtaagtcaa	atgaggaagt	gcagagggtt	tttttacata	tatatagcaa	2820
ccttgtcaag	tggtcctcac	aagagtcata	aatactttgt	aattagcaca	gtatattcag	2880
cagigtataa	ctctacaaat	agtaccttat	attagtgtag	tattatatca	atatcttatg	2940
tataattctt	atattaatac	cttatgcata	attggattca	aacattgaag	gtctatttta	3000
gtgttcttca	aaatgtgctt	ccctgaccta	ctgaaataga	aacttggtga	tgaagttcaa	3060
gaatttgtat	tctaatcatc	tcaaacaatt	cctaaagaca	ctgattttaa	aatatctagt	3120
ctaggcccca	ttgtgtaata	gttagcactc	taaaagatga	aaaagaaaat	agtctatgtg	3180
ccaaccactt	cattagtact	tatgaattta	aaaatgaaaa	agtctggtac	aggagacaag	3240
tatatatata	aaattataat	gcagtgtgat	aaatccatta	tagtatgtat	aagatacaga	3300
agagggactt	taaacttgag	aattcaatag	agataataaa	tgggtaggag	ggaaatagaa	3360
aactttggtg	ccacaaaagc	aaagtatgta	tggtattgcc	aataatagct	accatctatt	3420
gagtgcttta	ctacctgtca	ggtactgtat	tatataaact	ccattttaac	tgtacctcat	3480
tttgcagata	ctcaggcaca	aggaggtggt	tatttgtcca	aactggaacc	aagattcaaa	3540
cccagacaga	gtcttaagca	catttttaat	cactaactaa	cttgagatgc	ctaaatgcca	3600
aatactgttg	ggagttcaag	tggttcttga	ttagcaaaat	ctatttttat	tagtgcaaaa	3660
gaaacaccac	agcttataaa	gtattatgaa	ttcaataaat	ggagtcttaa	ctaatgagat	3720
attattttct	agaatggtgt	agctgagagt	atgtgtgatt	caactgaaag	gaataatgtt	3780
taatcagtga	ctcttactat	atacaggaaa	aggtgcagtt	ctgtctttca	aatctgcctc	3840
cttaccatat	tggcttacat	ccctcatgct	gttttcttgt	gtttgctaga	aagttgttgc	3900
caagccaaat	gtcatggcca	tgttgaaggc	aaggaag			3937

<211> 4720

<212> DNA

<213> Homo sapiens

ctcatgcttc	cataatagtt	ctgggataat	tctaaacaca	agccattttt	ctaaggagag	60
tccacattag	agaggtcttt	gttttgtatt	caagatgatc	aaaattatga	actgggaagt	120
tagtccctgg	ggtgtcctgg	ctggcctttg	gaaatcttca	ctacatcttt	ctgggttgga	180
atteteacea	cagcctgaac	gtggggctgt	atctgagctg	tctctgagtg	ctgtccattt	240
gatatatcga	gtactgggtg	tttaccaggg	ctcttcaagc	cactgggaga	aacagctaaa	300
gagtaaccta	ctgatttgag	atgtggattt	gtgccccatc	cctttctcct	tgtttcccac	360

aggagtttta	tctcaaactc	ctaagccatt	tttaaggaga	tcactggaac	aaactccaaa	420
cctaccctct	aatagtcaag	tttacctgaa	ttttttcagt	tctctcggga	gaagactaat	480
cacacattgt	agtaccaact	tggactcttc	atgtgctttt	cttaactgat	tagagttaac	540
acctcagcta	aagtgtatag	aacatacatg	gggcttcatc	aggcttcaga	atcagtttca	600
ctagatgtgc	tatgtaggag	gccacggaaa	aattactgta	gtagtaaaag	ttatcagttc	660
tgatgtaaac	aatcattttg	tcccatatta	taaataaatt	ggcctgaaaa	tatcttttca	720
talgtgagga	ataagtatat	gatgcctttc	tcctttaaag	tatgaactgc	taaaagacag	780
ggataacgtg	tattctgtat	tccagcagcc	acagtgtgtt	tctggtcttt	gtaccaggtg	840
ctcaggaagt	gttttcactg	gcttgggttg	actacttgcc	atctgctctc	tgagcattca	900
tttctgaatg	aaaggggaga	aagtgaaagg	agaggtggga	agaaagagga	agctgcagaa	960
atacgaggaa	acagctggag	gagggaggtg	aagttgagga	ggtaaggtca	gtaaaacaaa	1020
aagctagcag	agggcagggt	caggcccttg	gggtagaggg	ctaattaact	tctgtcagct	1080
agttgaatag	agccttgtgt	gctttgttag	agaccaaagg	tacttcaaag	gaaaaaaatc	1140
tagattcttc	cctgtgtacc	ttaataattg	ttcatcaggt	caaaatctat	cctgtcctct	1200
aggaattctg	gtcttccctc	aggcctagca	gagagctttc	igccactact	caggcaacca	1260
agggtgaagt	gcttcaagta	gtatttgtgg	acagcagcag	gtaagcttga	tgtgttattc	1320
acagcttaaa	gagtagatgc	tgagtacagc	tgttgtccat	gigtagagci	tttaataacc	1380
agcgcagcag	gcccttcac	ctgcttttat	gcctggacca	gatgactgaa	tgtagaactt	1440
taggcacttt	tttttttt	gagacggagt	ctcggtttgt	tgcccaggct	ggagtgcagt	1500
ggcgcaatct	cggctcactg	caagctctgc	cccccgggtt	cacgccattc	tcttgcctca	1560
gcctcccaag	tagctgggac	tacagactcc	caccaccatg	cccggctaat	ttttatattt	1620
tttagtagag	acagggtttc	accgtgttag	ccaggatggt	ctcaatctcc	tgacctggtg	1680
atccacctgc	cttggcctcc	caaagtgctg	ggattacagg	tgtgagccac	cagateggee	1740
ctttaggcac	tttctacttc	tcaaggtcaa	gaaacatcct	ttaaaaaagtt	aattcccttt	1800
tctggagcct	aagccagatc	ttatctaggc	cttgtgttgc	catctgttag	cattgatttc	1860
tggaatggag	cagctttctc	aaagtttggt	cttgctagtc	atgaggtcat	gtcagtgtct	1920
taggtcactg	ctgctcacct	tccttaccca	gggagtatac	tgcataggtt	tctgaacacc	1980
tgttttcatt	attcactgtt	cctctcactg	ccaagaatgg	agggaccctc	agttgaagat	2040
caaattgact	ctgaagaaaa	actggagatg	tttctcttgg	agtttggata	gagtattcac	2100
ttgataacat	gtttttcccc	tgccttgctc	ttcacaagaa	catciggcca	ggcattaaca	2160
attagtaaat	tttttgcat	atgaacagta	tttttctggt	catglagatg	ggtgcacatg	2220
acactaaaca	gcattgttta	gtgttatccc	tcttaactgg	tgggttgtat	ttggggtgga	2280
ggctgtagcc	gaggagaaga	cattcacctc	tgtactcgag	aaactttgtg	taggaattta	2340
gtttatttt	ttatttttt	aatttttat	tttttactac	ttttactgtt	agcacaatgc	2400
tataattgag	ctaatctttg	tagtttggtg	caggaccacc	aagtttgtgt	gacccattac	2460

```
ctactttttc catgctcagc cattaccctg tcctggggca tctgagggca glaaggaaca
                                                                    2520
ggtgtccaaa ggaggaatgt tggtgcctat gagtatgttt tccagttgta ttgaatttct
                                                                    2580
tacttggtgt atttttgact tgtcttagtt tctttccttg tggtctatgc tattttactt
                                                                    2640
gcgattigtt ggatattete eetgleatta aagagitgia aaaiggaagi tagiitetet
                                                                    2700
atgcaaatgc tttaatggat gaagctgata ggtttagcat tgatttttgc tggtgtcctt
                                                                    2760
caacaagcat gaaggtgata aatgtgtttc catggcttta gactcatttt tgaagtcttg
                                                                    2820
galigigiga acalicitag aaacaalaaa algilitaat taaaagccci cgaclaccag
                                                                    2880
ctgaattcag tgtctactag gaaaatgggt agatttgtta cattgtccct ttgctctcta
                                                                    2940
tgactttgtt ccagttgtca aggaacttaa atgggtattc aggaaaaaga attcttgttt
                                                                    3000
ccelttecte accttgecag ttaaataact eetggtgaca etteaggtgg tagaattgaa
                                                                    3060
acacaaacct gacttetgae cacatgggte aaaggeaaaa ggeaaatgge tteaaageee
                                                                    3120
ttagtgtget tatecagtte aggeagtgag gagataacet etgettteet eeetgaggag
                                                                    3180
ttiggagiat ttaaggggggg algggggggg tgicactiig aaaataigii gciititcic
                                                                    3240
ctgattgtat tgaggctgat atggaagggt tatttctttc tggccaatac tttttggtat
                                                                    3300
ttctaaatat tgcaatcttg atttttacta ttaaatttgt taattgtcag ttctggcttt
                                                                    3360
ttigcataaa gagitggicc altaacitgc caattigaag citciaacia gataticcci
                                                                    3420
actgaaagtt ttggatttgt ttttagtttg tggagcagtc ttagctgggg acaggtaatt
                                                                    3480
gacaacggca gagatacttt cttttcctag gattctaagt ctgtaatcca catcctcaat
                                                                    3540
gtattcacag gactttaaaa ttctctccaa atgaggaagg aaatatcctg ttgctttcta
                                                                    3600
atgtttacta aaagttgtgt ttagaacaac agattttaat aggcatcttc ctttgttatg
                                                                    3660
                                                                    3720
tgtcattagc cctttgcccg tttaccttag ggctctttga aggagaaatg gatgtgagaa
aacctgtcac ttggcgaaag taaaagggat aattaactgg ctcagagctt atgtgcagag
                                                                    3780
ttecaageee caaagttaat etagaaceae tegataacae caataaaaat atttatttea
                                                                    3840
catetgitat atatetggaa aatgitetaa geatettaea eatatitete attaaateea
                                                                    3900
caggigacca tigigaggia galattiigi telaattiic cagaigagga ageigagacc
                                                                    3960
ctaaaagget gaceggttee etgatgtgt acetgettet getaetgate caaactgeag
                                                                    4020
aacticteat teateceeaa ggeeteeagg eagtateeaa tggggaatea getetaaaag
                                                                    4080
gaaccagacc aacgttttcc agccccttca ttctggtgac tgaggggagg aaagaatggg
                                                                   4140
agggggtatt cttgtctagt ggatggaaag gaaacacact gtcaaattac tatatctcct
                                                                    4200
lgglitteta ttacagtaga altetecage catatitita itgiciatgg gggaagiigg
                                                                   4260
agatggtgac ctigattaga agtgtctgga gggggataaa tggaggggat aagattcagt
                                                                   4320
tggttttgga aaatgitaaa gictiaaaat aatgegteea tetgaagaat tiittetaaa
                                                                   4380
accagagtit ataaaaatat cactgataca geetgeeeee teatticeet geeacaggag
                                                                   4440
atgictigga ciagagacac tigittaata atagcitgic icigalatic ccagiagcit
                                                                    4500
ccclctgtgt gaggaaagga tagaaatgii caggacatca tcatacaggc iccicaicta
                                                                   4560
caaagtteea gtageagtga egeetaeaeg gaagaettgg aactgeaaac aggetggggt
                                                                   4620
```

cacctcagtg acatctgacg ctgtccaacc agaagttcga tttttgttct gggggtgaag 4680 gaggaaacag actgtactaa aggactaaaa taatttgtct 4720

<210> 2024

<211> 3531

<212> DNA

<213> Homo sapiens

<400> 2024

60 agaataaagc tttcagcaag tttggatctt tttctgccac cttagaaaat ggaatctgcc tctcgataag ttactatgga tcaaatggaa tggcaccaga agataaggat cctgatttag 120 aaacaatatt gaatateeet teageactea eteeaacagt ggtteetgtt atagtgaeeg 180 240 tteeteaaag caaagetaaa gggaaaataa aaggeaaaga aaaacecaaa gaateeetta 300 aagaagaaga acacccaaaa gaagaagaga aaaaggaaga agaagtagaa ccagaacctg ttttacaaga gactttggat gttcccacct tccagagcct aaatgtgtct tgccccagtg 360 ggctcctgtt gactttcatt ggacaagaat ctacaggtca atatgttata gatgaggaac 420 480 ccacctggga catcatggtc cgtcagagct accccagag ggtgaagcac tatgagttct 540 ataaaacggt gatgccaccc gcagagcagg aggcttcaag ggttatcacc agtcaaggca 600 ctgttgtcaa atatatgttg gatggatcca cacagattct ctttgcagat ggtgctgtga geaggagtee caatteaggt cttatttgte etecttetga aatgeeagea aegeeteaca 660 gtggagattt gatggactct attictcagc agaaatcaga aacgatacca tctgagatta 720 780 ccaacacaaa gaaaggaaaa agtcacaaaa gtcagtcatc aatggcccat aagggtgaaa 840 tecatgacec tectecagag geagtteaaa eigtaaciee tgiggaggti cacataggea 900 cetggtttac aaccacact gaaggaaatc ggatcggcac caaaggatta gaaagaatag 960 cagacttgac cccattgtta tectttcagg ccacagatec tgtcaatgga acggttatga caactcgaga agacaaagtt gtcatagttg aaaggaaaga tggtactcgg atagtggatc 1020 1080 atgctgatgg taccagaatc acaacctttt atcaagttta tgaagatcaa attattctgc cagatgatca agaaacaacc gagggtcctc ggactgtcac caggcaggtg aagtgtatgc 1140 gggtagaaag clcacgclat gccactgtla tcgccaactg tgaggacagt agctgctgtg 1200 1260 ccacctttgg agatggaaca actattattg caaagccaca gggaacatac caggtgttac ctccaaacac aggctctctt tataltgaca aggattgttc agctgtgtac tgccatgagt 1320 1380 caagcagtaa tatatactat ccttttcaaa agcgtgagca gctgcgagct ggcaggtaca teatgaggea tacticagag gitatetgig aggitetgga teetgaggga aacaetilite 1440 aggicatggc tgalggtagc atatcaacta tattacciga aaaaaaatig gaagacgati 1500 taaatgagaa aactgagggc tatgatagtc tgtcctctat gcaccttgaa aagaatcatc 1560

```
agcaaatcta tggtgaacat gtccccaggt tttttgttat gtatgctgat ggatcaggaa
                                                                    1620
                                                                    1680
tggaacttct tcgagacagt gacatagaag aatatctatc tttggcatat aaagaatcaa
                                                                    1740
atactgttgt tctccaagag ccagtgcagg aacagccagg caccctaacc atcacagtcc
                                                                    1800
ttegecettt ecatgaagea teaceatgge aagtaaaaaa ggaagataca attgteeete
ctaatctccg gtcaaggtca tgggaaacat ttccctcagt tgagaaaaaa actccaggac
                                                                    1860
                                                                    1920
ctccgtttgg tactcagatt tggaaaggcc tttgcattga gtccaaacag ctagtgagtg
                                                                    1980
cccgggtgc catactcaag agccccagtg tgctacagat gcgccaattc attcagcatg
                                                                    2040
aggtcataaa gaatgaggtg aaactgaggc tgcaggtttc ccttaaggat tacataaact
                                                                    2100
atattctaaa gaaagaagat gagctgcagg aaatgatggc taaagattcc agaactgagg
                                                                    2160
aggagagag caatgctgct gatctcctca agctggttat gtctttccct aaaatggagg
                                                                    2220
aaactacaaa aagtcatgtt actgaagttg cagctcacct aactgattta ttcaagcagt
ctttggctac gcctccaaaa tgcccaccag acacatttgg taaagatttc tttgaaaaga
                                                                    2280
                                                                    2340
catggagaca cacagcatcc tcaaaacgct ggaaagaaaa gatagacaaa acgaggaagg
aaattgagac aacacagaat tacctaatgg atattaagaa ccgcataata ccaccctttt
                                                                    2400
                                                                    2460
ttaaatctga attgaaccag ttatatcagt ctcagtataa tcacctggac agtctttcca
aaaaactgcc ttcttttaca aagaaaaatg aagatgcaaa cgaaacagct gttcaagata
                                                                    2520
catctgatct taatctagat ttcaagccac ataaggtttc agaacagaaa tcctcaggtg
                                                                    2580
tgcctagtct tccaaaacca gagatttctg cagataagaa ggatttcact gctcagaacc
                                                                    2640
                                                                    2700
aaactgaaaa tttaacaaaa tctcctgaag aagcagaatc ttatgagccc gtgaaaattc
                                                                    2760
caacccagtc cttgctgcag gatgttgcgg gacaaacaag aaaagaaaaa gtgaagttgc
                                                                    2820
ctcattattt gctgagttcc aagcctaagt ctcaacctct tgcaaaggtg caagattctg
tiggaggaaa agigaacaca teeteigiig cateigeige cattaataat geaaagicat
                                                                    2880
                                                                    2940
ccctttttgg gttccatctt ctcccatcat cagtcaagtt tggagtgctt aaggaaggac
atacctatgc cacagttgta aagetcaaga atgttggagt ggacttetgc aggtttaaag
                                                                    3000
                                                                    3060
taaagcagcc cccacccagc acaggactga aagtgactta caaacctgga cctgtggcag
                                                                    3120
ctggtatgca gacagaactg aatatagagt tatttgccac agctgttgga gaggatgggg
                                                                    3180
ccaagggatc agcacacatc tctcacaata tcgagattat gacagagcat gaggttctgt
tectacetgt ggaagcaact gttttaacaa geagcaatta tgataaacga eeaaaagact
                                                                    3240
                                                                    3300
ttccccaggg aaaagaaaat ccaatggtcc agagaacttc tacaatttat tcctccacac
ttggagtett catgietegt aaagtitete cacattaggi acattiette teggiacaac
                                                                    3360
                                                                    3420
teaatageet eeataateet eteageetae agaggatgag aaaggaaaga agteateaca
acatacteca teateceagg acactgaaac tggaagaact gaccagaaat ttgccaaatg
                                                                    3480
aaatagette aatetgitta ataaagaegi gegaatagag igeeaaaaag e
                                                                    3531
```

```
<211> 3361
```

<213> Homo sapiens

```
agctctggga gaggagcccc agccgtgaga ttcccaggag tttccacttg gtgaccagca
                                                                      60
ctgaacaca accaccaacc atggagtttg ggcttagctg ggttttcctt gttgctattt
                                                                     120
taaaaggtgt ccaatgtgag gtgcagctgg tggagtcggg gggagccttg gtgcagccag
                                                                     180
ggcggtccct gagactctcc tgtaaatctt ctggattcac ttttggtgat tatggtatca
                                                                     240
gttgggtccg ccaggctcca ggaaaggggc tggagtgggt aggtttcatt agaaacaaag
                                                                     300
cttttggtgg gacaacaata tacgccgcgt ctgtggaagg cagattctcc atctcaagag
                                                                     360
atgatteeaa aggegtegee tatetgeaaa tgageageet geaaacegag gacacageeg
                                                                     420
tatactactg tactagagac atcittgtta ctgggatcta tcattactac tttgactact
                                                                     480
ggggccaggg aaccetggte accgteteet caggtgagte etcacaacet etcteetget
                                                                     540
ttcagtctga aggttttcac tacatttttg ggggcaaata tgtgtgctgg gtctcctgcc
                                                                     600
aaaagageeg eggaacagtg gggggggete gggaaaatgt eetgaggeag eggeggeeaa
                                                                     660
acagacgagt gccaagggct ccagatgttc cttcctcttc agcccaacag cacgggtctg
                                                                     720
tetgtggcca gggccaccet gggcctctgg ggtcccatgc ccaacaaccc ccgggcctc
                                                                     780
cccgggttca gtctgagagg gtcccaggga cggagcgggg cgccagttct tgcctgaggt
                                                                     840
cctgacattg ttctcacaat gtgacaactg cttcgacccc tgggggccagg gaaccctggt
                                                                     900
caccgtetee teaggtgagt ceteaceace ecetetetga gtecaettag egagaeteag
                                                                     960
cttgccaggg tctcagggtc agagtcttgg aggcattttg gaggtcagga aagaaacctg
                                                                    1020
gggagaggga cccttcgaaa gggaacccag cctgtcctcc ccaagtccgg ccacagatgt
                                                                    1080
eggeageigg ggggeieett eggeiggigi ggggigaeet eteleegeit eaceiggege
                                                                    1140
attctcaggg gctgtcgtgg tgattgcgtg gtgggactct gtcccgctcc aaggcacccg
                                                                    1200
ctctctggga cgggtgcccc cccggggttt ttggactcct gggggtgact ttacagccgt
                                                                    1260
ctgcttgcag ttggacttcc caggtcgaca gtggtctggc ttctgagggg tcaggccaga
                                                                    1320
atatgggaca aaccaggggt cttagtgatg gctgaggaat gtgtctcagg agcggtgtct
                                                                    1380
gtaggactgt aagatcgctg cacagcagcg aatcgtggaa tatcttcttt agaattatga
                                                                    1440
ggigcgcigi gigicaacci gcaictiaaa ticitiatig gciggaaaga gaacigicgg
                                                                    1500
agtgggtgat tecagecagg agggacgegt ageceeggte ttgatgagag cagggttggg
                                                                    1560
ggcaggggta gcccagaaac ggtggctgcc gtcctgacag gggcttaggg aggctccagg
                                                                    1620
acctcagtgc cttgaagctg gtttccatga gaaaaggatt gtttatctta ggaggcatgc
                                                                    1680
ttactgttaa aagacaggat atgtttgaag tggcttctga gaaaaatggt taagaaaatt
                                                                    1740
atgacttaaa aalgigagag attitcaagi ctattaatti tiltaacigi ccaagialit
                                                                    1800
gaaattetta teattigatt aacacceatg agtgatatgt gtetggaatt gaggeeaaag
                                                                    1860
```

```
1920
caagctcagc taagaaatac tagcacagtg ctgtcggccc cgatgcggga ctgcgttttg
                                                                    1980
accatcataa atcaagttta tittittaat taattgagcg aagciggaag cagatgaiga
                                                                    2040
attagagtca agatggctgc atgggggtct ccggcaccca cagcaggtgg caggaagcag
                                                                    2100
gtcaccgcga gagtctatti taggaagcaa aaaaacacaa tiggiaaatt tatcactict
ggttgtgaag aggtggtttt gcccaggccc agatctgaaa gtgctctact gagcaaaaca
                                                                    2160
                                                                    2220
acacctggac aatttgcgtt tctaaaataa ggcgaggctg accgaaactg aaaaggcttt
                                                                    2280
ttttaactat ctgaatttca tttccaatct tagcttatca actgctagtt tgtgcaaaca
                                                                    2340
gcatatcaac ttctaaactg cattcatttt taaagtaaga tgtttaagaa attaaacagt
                                                                    2400
cttagggaga gtttatgact gtattcaaaa agtttttaa attagcttgt tatcccttca
tgtgataact aatctcaaat actttttcga tacctcagag cattattttc ataatgactg
                                                                    2460
                                                                    2520
tgttcacaat ctttttaggt taactcgttt tctctttgtg attaaggaga aacactttga
tattetgata gagtggeett eattttagta ttttteaaga eeacttttea aetaeteaet
                                                                    2580
                                                                    2640
ttaggacaag ttttaggtaa aatgtgcatc attatcctga attatttcag ttaagcatgt
tagtiggigg cataagagaa aactcaatca gatagtigcig aagacaggac tigtiggagaca
                                                                    2700
                                                                    2760
ccttagaagg acagattetg ttccgaatca ccgatgcggc gtcagcagga ctggcctagc
ggaggetetg ggagggtgge tgecaggeec ggeetggget ttgggtetec eeggaetace
                                                                    2820
cagagetggg atgegtgget tetgetgeeg ggeegaetgg etgegeagge eccagecett
                                                                    2880
gttagtggac ttggaggaat gattccatgc caaagctttg caaggctcgc agtgaccagg
                                                                    2940
                                                                    3000
cgcccgacat ggtgagagac aggcagccgc cgctgctgca tttgcttctc ttaaaacttt
                                                                    3060
gtatttgacg tcttatttcc actagaaggg gaactggtct taattgcttg atgaagagca
                                                                    3120
ggagactcat ttatgtgagt cttttgagtg accattgtct gggtcactcc catttaactt
tecetaaage eeatttgaag gagaggtege aegagetget eeacaacete tgaatgggga
                                                                    3180
                                                                    3240
tggcatgggt aatgatgett gagaacatac caageeccac tggcategee ettgtetaag
teattgactg taggteatea tegeaceett gaaagtagee eatgeettee aaagegattt
                                                                    3300
                                                                    3360
atggtaaatg gcagaatttt aagtggcaaa ttcagataaa atgcatttct tggttgtttc
                                                                    3361
```

<211> 3527

<212> DNA

<213> Homo sapiens

<400> 2026

cttttctcta ttaggaagta ccaccaagaa cagggaagga caagccagag gctggaggaa 60 gatacctgca gaacacagac ctgacaaagg atcagtatca aaacatataa gaattttgac 120

aaatgaataa	aaagagtaca	aataacccaa	cataaagtca	aaaggcgtga	tcaggcattt	180
cacagaagca	aacacctttg	gtggatgccc	atgaggagag	gcgcagtcac	atcagtgccc	240
aggagatgca	aacccagatc	ccaggggtgt	gcatcccacc	cgttctgcct	gtaggatctg	300
caaacccggc	aaaacctagt	tctagagaga	ctggattcac	tgcatgtctt	catcactgct	360
ggagggagcg	cagactgcta	tcgcctctta	gaaaatgact	tagttctcat	gtaatttggg	420
cattcacaca	tcctcatcct	agatccagct	tttccactcc	cgcacacgta	ctggaaaacc	480
tgtacaggaa	catccactgc	agcactgctc	ataccaaaca	caacctacat	gttctctgca	540
cagagagggg	agaagagccg	gtcagttcac	tcagtggact	ctgtgctcaa	tagtaggtgt	600
gaataagccg	cagccgccca	gaccgcatgg	gccaacctca	gtccgagaat	gcggagtgaa	660
aacacaggtc	taacgatcac	acatggcaag	ataattatct	tgcaaagaaa	actcacttat	720
tgttctgcca	tacatatatg	taccataaaa	tcactccccg	caccttccac	tctgaaaaaa	780
caaaggaatt	ctaggcacaa	agttcaggat	catggttaac	tgaggggaga	gaacagggag	840
tatgatggca	agatagaagg	tatcgttcac	atccaagttt	ataggttggg	ttcttgattt	900
agtcattatt	caaaggctaa	taactaaata	aaaggtagct	agcgtgagag	tgcaacatga	960
accaaagatc	atgactggct	ttgcgcatcg	aggggccatt	aaagagtcta	cttttcatgt	1020
tatcacttaa	aatcattttg	cacccaccag	ggcatgagca	tctcgtgctg	gcaaacacca	1080
catgaccgtg	gtgacctcag	ggccagcccg	ggggtcatct	tgaatctctc	ctgctgaaga	1140
gacccaggag	ggtaacacac	gcccctccaa	tctctgagtt	ctaggaaatg	aacacctggt	1200
atttaaaggg	gctgacataa	tgcaaatcat	ctgatgaaat	gtttgtttta	gttcacttaa	1260
agatcaacac	gagagtcttc	actctgaatg	ggccacacct	gaattaagag	aatccttcac	1320
tctctgcgtc	ggatgcacaa	accagtcctc	ctggtgctca	caggggctag	cagcaagtcc	1380
agaccttgta	tggtgagggc	ggggggggat	ggtgaactta	gggttcagcg	aaaccgccac	1440
ttgcaaacac	accccaccgc	aggtgccctt	gatgtgtaca	cacgtccttg	agaagctggg	1500
ggcaaggcct	tgcgggtgag	accacgctca	gcagctcaca	cctttaccaa	gtactaggac	1560
ttctttgggg	ttgggttgag	gggtgatccc	aatctgagtc	tatggtatga	ctcaggggag	1620
aacaggtcac	cgggtgctag	gagagctgtc	catagaggac	acagccccaa	aggattagaa	1680
ccaggagaaa	ggtagagtct	gactcagggt	gaggaacaca	catatattgg	tgctgcccga	1740
aggggaactg	cctcgtgagc	gtctgggaac	tcttactgca	ggtgctcagc	agatgcttgg	1800
tgccctgcag	ggacgtgctg	gcctcgatcc	tcgcgaggca	gagccccgga	ctaggagaca	1860
gttcaggtcc	tgcataacct	gagtgtccac	agggcccagc	tagtcctcaa	gctggggctc	1920
gcccagtggc	tgctccctct	gcttctccca	tcctgactcc	gcctgctcct	ctttggagaa	1980
gtgaggggtg	aggggcccag	aggcaggggc	tggggtgggc	tctgctgcat	gtggaggcga	2040
aggaggagag	gggaggggag	gcagcatcaa	agccagtctc	tctagctcag	actctgggtg	2100
gtttgggtgg	gtcctgcccc	ctggcctgtt	cccgtctgtg	gggtcccact	gcttgggtgg	2160
tgtagcttca	ccccatettc	ccacaccggg	gtgcctggtg	ctcagcctcc	cctcaggtag	2220
gctctgtgcc	tcctgattcc	tcaccgtggg	tggtccctcc	tgcctgcagc	ctctaaggcc	2280

cctgagagca	gtcagtcagt	cccaaagtcc	ccaccagcgc	tgctgactca	cttccgatgt	2340
ccttgctgcc	gtgttcaggg	agctggaggg	ccaggctgac	ccgcttgggg	gcttcctcca	2400
tgttctcgag	ctgcgccgag	gctgtgggtt	ctaggagaag	ccaggcggtg	accacacggc	2460
gcagctgctt	tgcacccggg	atggtcctgg	ggccaccctt	ttgagtgctt	ctatatctca	2520
gggagcacgg	atgtccctgg	tggggaccag	gctccctgcg	tggccccagc	acctgtcggc	2580
cccagagctg	cctccctga	${\tt agggctggcc}$	tcaccctcct	gctgaccctc	tggaggggct	2640
cggccttccc	cttgcagggc	ccctcagag	ctgcttcagg	gacagccacc	actgatcatg	2700
ctgagaggcc	ccatcctcac	ggctgatgcg	gttgctttct	tcttagggtc	aaattctgca	2760
ttcctctcct	tccacccctg	cttcttggag	gctgtggcac	ccctgctct	ttctgagctg	2820
ccctcagtct	gtactgacct	tcctcatgcc	ctcgccccca	actgcatcac	ttcttatgca	2880
gggatctcaa	ccgcaccctc	gggcacttca	tatccgcttc	catagctgca	agtacaacgg	2940
gccccctct	gtactccaga	tctcacctgc	ccaccactgg	gcatcccggg	cagctgcctg	3000
ccctctcctc	agacaccttg	ctgggggctc	tctccctgc	tcaccgtgcg	gcagggaccc	3060
cagggctctg	gtccctgctt	gccactctcc	ttgctgcatt	tccctccctc	ctctgcctga	3120
ggagttttg	ctcagagcgt	gttcattaaa	ctggtgacta	ggctctgttg	gggagttcca	3180
tgaggatgac	cacctggcct	tccaggtgag	aggcaagggc	cagagaggtc	ccctctgggg	3240
cagggtcgcg	cctgcctcac	tcctgccaac	atgtctcagg	gcttctgtgt	cagaatcaca	3300
ggcagattcc	cagagcggca	ctcacccagt	aaacccggtg	ggaagggccc	aaggcacctg	3360
ggcccatcag	ccttgctgcc	accgggaaga	tcttgccagg	acagtggcgg	aggatttgcc	3420
ggaccacact	cggagtggcg	ggttagaccc	tcatggcctc	ctgcccatgg	tttactaaaa	3480
caaagctcag	agccctactt	tggcaaataa	agctgctgta	atgtctc		3527

<211> 3677

<212> DNA

<213≻ Homo sapiens

taticititg	aagagtgcat	ttcaagctgc	caaggtggag	agagggatta	cagaaaggag	60
aacaccttat	ttcaggaacg	tggatggagc	tggtggccat	tattcttagt	aaactcatgc	120
aggaacaaaa	ccagatactg	catgttccca	cttacaggtg	agagctaaat	gttgagaaca	180
catggacaca	cagagaggaa	caacagacac	tggtgcctac	ttgaggatgg	aaggtggaag	240
gagggagatg	agcagaaaaa	ataactgttg	gcttagtacc	tgagtcacaa	aatggaacag	300
ctggaactga	gtcttcaaaa	agctgttcaa	gagttctgaa	gaccagctgc	ttctggatga	360
tggaatgtac	agtaagacca	gaacgtagat	ggagctggtg	gccattattc	ttagtaaact	420

catgcaggaa	cagaaaaacca	aatactgcac	gttcccactt	acagcacaag	aaccatcaat	480
cagcacagaa	gacttctgtg	accccaaact	atgtggggat	ttctccctag	caacaagcaa	540
gcaatcagtt	tggcaacaaa	cactgactgg	atgtcttcca	attcaattcc	aacactatct	600
acctggaaat	agtgtctgat	cccacaggaa	acggggtttc	accatattat	ccaggctggt	660
cttgaactcc	tgggctgaag	aaatccacac	acctcgtcct	cccaaagtgt	tgggattaca	720
ggcgtgaacc	accacacccc	agttcaatcc	attctacaca	aactgtaagg	atagtttttc.	780
taaaacagta	catggatcaa	ttaaaactga	tcagctttac	actaaaactg	atcagagttc	840
tgatcagttc	aagaactttg	tctgtgtagt	caagtcccaa	atcatcattc	tggcacttaa	900
ggaagaatgg	ccacagcctc	atctccctat	tattcccttt	atgaaagctt	atggtatctt	960
tggtttagca	ctgattccac	ccttccctgc	cttttatata	attggtgtat	ttatcttttg	1020
tctagattgt	gagatccttg	gaaacaggta	ttatgcatat	aaactcaata	actttatttc	1080
tttttctgcc	aagtgaacaa	agacctccaa	attgtagtca	catgtaatac	agaacactgg	1140
tatiggicat	atctccatct	ctgatccacc	cttcctccca	tgcaagctat	agatgcatat	1200
catctatttt	gaattgccta	ttgaaatgtc	cctttgaata	tctaataggc	atttcaagtt	1260
taatatattc	agaatagttt	ttcctccatt	gtttactacc	tgccagttgc	ttaaggccaa	1320
aatctatgaa	tcattcttga	ttcttctctc	attttccata	ttcaatccat	cagaagtttt	1380
aatggcttta	tctccaaaat	atatcttaac	aatggagcaa	aatttgagta	ccatagaacc	1440
agattaggga	taactgtagg	aatgaaggag	tgatacacct	aggaatagga	aagtagtctg	1500
aagccagatg	ataaaggtcc	ttttatgcca	aactaagaag	ttcagatatt	attgttagag	1560
gagagctatt	gaggttttg	agtagggcac	ttatatattc	attttgtact	tcaggaatga	1620
atcagtagag	gggataaatg	agggtaccct	caaaattgca	agcaatgaag	tattagaact	1680
gaattttag	ggacagcaat	accagectaa	ctggctgcat	aagagaagac	tgggtgtttg	1740
ggagaggcaa	aaaataaggc	caacagaaaa	ggtagggtag	ggacagaata	taaagggctt	1800
gaaagccagg	taaaagcata	taaactggga	tgcctgggtt	ctaataactg	tttccctact	1860
tggagaaact	ccccttatct	tttttaaacc	ccagtttcct	tttcctggga	tttgcttcag	1920
ctataaatgt	tgtaattttc	tataatgctc	tgacctgcta	cagtggctct	gaaaccttga	1980
ctgcacactg	gaatcaccta	gagagcttta	aaagctactg	atggctagat	ctcactacca	2040
aagattcaga	tttatctggt	cttaggtgca	gcctggacac	tgagatattt	aaaagttctc	2100
caggtgattc	taatgtgcag	ccaaggttga	gatcaactca	tgtagaaaat	agtgaagcac	2160
taagattett	aagcatggta	ataatatgtt	aaaatttagt	ttagtttttt	tgttttttt	2220
gttttttcca	agacggagtc	ttgctctgtc	gcccaggctg	gagtacaatg	gcatgatttc	2280
aactcactgc	aacctccacc	tcctgggttc	aagcgattct	cctgcctcag	cctcccgagt	2340
agctaggatt	acaggtgcgc	accacaacgt	ctggccaatt	ttttgtatit	ttagtagaga	2400
cagggtttca	ccatgttggc	cagggtggtc	tataactcct	gaccttgtaa	tctgcctgcc	2460
tcagcctccc	aaagtgctgg	gattacaggc	atgagccacc	gcgcctggct	gaaaaaaggt	2520

attttaagaa	agactaacag	gaatatacag	actagtaggg	aaagactaca	gaagatcaac	2580
tagaattttg	caataatcca	ggagaaaagt	ttagtaaggg	ctggattagc	atacatgcaa	2640
tgatgttgta	gggaggaaga	tgaatgcaag	aaacatttgg	agagaaggag	caccaggatt	2700
cagtaagtga	ttgaatgtta	aatctgagca	aaaggaaaaa	aaatatggtc	aagtttctag	2760
catagaagaa	taatagactc	cttaacaaaa	ttaaagtagt	tgtgaaacag	ctggttaatc	2820
aatattattg	agaatatgga	aactaacatt	aaattctaag	tcggggtcta	acctacgtgc	2880
cttacataca	ttatctcatt	taatctttac	aaccaccata	taaatactac	tatcattccg	2940
attttacagt	tttagaaact	gagtaagagt	aattaaatta	tttgcccaaa	gttacacagt	3000
aaatggtaga	gaagacattt	gacctcaact	ggtctaacta	cttttcctca	taggaagatg	3060
accagtttac	atatggaatc	tgttgaattt	gagcaaacaa	ctcaaaaaaag	caaaatggct	3120
atagaggcca	gatgggaaca	taaatgagtg	aatcaagtca	gatgcaactg	tggagaaatc	3180
aaaacatcga	gagaaggtag	ttctacttag	ttatgcttga	atgttgccct	atgagaattt	3240
caggcccagt	attgccatat	tttaagattt	ttcatgaaaa	gatggaaatc	tggatttgta	3300
tgcaaaaatt	tgtgtgaata	tcaaattcaa	gtgtttaaaa	ctactgtggc	tcaaactatg	3360
gcttcaagtt	tgcatctctg	agcaaaaggc	tgttggaaat	tcagaactgg	atgtaaagtg	3420
agagatctgg	gctgaaggta	aatgattagg	gaattcataa	gcacagagag	gatggtagat	3480
gcttccaaaa	cagtatgtgt	tagaatagta	accagcactt	gacatgatta	gttaaaataa	3540
ggcaaaaata	tatgagttaa	caagttagtc	aggacttaga	gaaaactgat	aaaactagca	3600
gtggaaaact	agcagactta	agtgggtata	tttaaaattc	aattttcaat	gaactaaaag	3660
ctaaattcca	gacaatg					3677

<211> 4143

<212> DNA

<213> Homo sapiens

aaaaatatgt	agaagatgaa	atggcaaggc	tccctgatag	attgtcagta	acttggcctg	60
aaggagatga	attattgcct	aatgagatta	ggcctgctgg	aacccctatt	ggtgcgttaa	120
gaattgaaat	actgaataaa	aaaggggaag	caatgcaaaa	gcttccagga	acaagccatg	180
gagggtcaaa	gaaactcctg	gttgagctca	aagttattt	acattettea	agtggaaata	240
aagagattat	ticgcatatt	agtcaacatg	gaggaaaatg	gccttactgg	tttaaaaaaaa	300
tggaaaatat	tcagaagttg	gggaattata	ccttgaaatt	acaagttgtg	ttgaatgaaa	360
gtaatgcaga	cacttatgca	ggaagaccac	taccatctaa	agcaattaag	ttttctgtta	420
aagtggttla	tctttacatt	atgaagaaat	aaccaaagga	ccaaattgtg	taattcgagg	480

tgttacagcc	aagggccctg	taaactcttg	tcaaggcaag	aattataatc	tgaaggttac	540
tctgcctggc	ttaaaagaag	actcacagat	tttgaaaatt	agattactac	ctggtcaccc	600
tcgtcgactg	aaagtgaaac	ctgattctga	aattttagtt	atagaaaatg	gaacagcttt	660
cccatttcag	gtggaagttt	tagatgaatc	agacaacata	acagcacaac	caaaattgat	720
tgttcattgt	aagttttcag	gtgctccaaa	ccttccagtc	tatgttgtag	attgcagtag	780
ttctggaacc	agtatttaa	caggatetge	aattcaagtt	cagaatatta	aaaaagacca	840
gacgcttaaa	gcaagaattg	aaatacctag	ttgtaaagat	gtggcacctg	tggagaagac	900
tattaagttg	cttcccagta	gccatgttgc	aagactacaa	atattcagtg	tagaaggaca	960
aaaggcaatt	cagatcaaac	atcaggatga	ggttaattgg	atagcgggtg	atattatgca	1020
taatcttatt	tttcaaatgt	atgatgaagg	agaaagagaa	atcaatataa	catcagcttt	1080
agcagaaaaa	attaaagtta	attggactcc	tgagattaac	aaagaacact	tgctacaggg	1140
tetgetteet	gatgtgcaag	taccaacatc	tgtaaaagat	atgcgctatt	gccaggtttc	1200
attccaagat	gatcatgtgt	ctttggaaag	tgcgtttaca	gtaagaccac	ttcctgatga	1260
acctaaacat	ttaaaatgtg	aaatgaaagg	aggaaaaaaca	gtacagatgg	gccaagagct	1320
tcaaggagaa	gtagttataa	taattacgga	tcagtacgga	aatcagattc	aagcattttc	1380
accaagttct	ttatcttctt	tgtcaattgc	tggggttgga	cttgatagct	caaatttgaa	1440
aacaaccttt	caggaaaaaca	cacagagtat	aagtgtaaga	ggcatcaaat	ttattccagg	1500
tcctcctgga	aataaggatc	tttgttttac	ttggcgtgag	ttttctgact	ttattcgagt	1560
gcaactaatt	tctggacctc	ctgctaaact	tctccttata	gactggccag	aactaaagga	1620
gtccattcca	gtgattaatg	gaagagattt	acagaaccct	attattgttc	aactttgtga	1680
tcagtgggat	aatccagcac	cggtacaaca	tgttaaaata	agtcttacaa	aagctagcaa	1740
tttaaagctc	atgccttcaa	accaacagca	taaaacagat	gagaaaggca	gggctaattt	1800
gggagtattc	agtgtttttg	cccctagggg	agagcatact	cttcaggtta	aagccatcta	1860
taacaaaagt	atcatagaag	gacctataat	taagttaatg	attcttccag	acccagaaaa	1920
acccgttcgt	ctcaatgtta	aatatgacaa	agatgcgtcc	ttcttagcag	ggggtctttt	1980
cactgatttt	atgattagtg	ttatttctga	agatgacagt	atcattaaaa	acattaatcc	2040
agcacgtatt	tccatgaaaa	tgtggaagct	gtctaccagt	gggaaccgac	ccccagcaaa	2100
tgcagaaaca	tttagttgta	ataaaataaa	agataatgac	aaagaagatg	gctgcttcta	2160
tttcagggat	aaagtaattc	ctaataaagt	ggggacatat	tgtatccagt	ttggttttat	2220
gatggataaa	acaaatattc	tcaacagtga	acaggttata	gttgaagtcc	tgcctaatca	2280
acctgtgaag	ttagtaccta	aaattaaacc	acctacacca	gctgtttcaa	atgttcgctc	2340
agttgccagt	aggaccttgg	tcagagatct	acatcttagt	atcacggatg	actacgacaa	2400
ccatactgga	attgatttgg	ttggcactat	aatagccacc	attaaaggct	ctaatgagga	2460
agatactgat	accccacttt	ttattgggaa	agttagaaca	cttgaattcc	ccttcgtgaa	2520
tggttcggct	gaaatcatga	gtctggtgct	ggcagaaagt	agtcctggaa	gggatagtac	2580
tgaatatttt	attgtatttg	agccccggct	accactttta	tcaagaacct	tagaaccata	2640

```
tatcctaccg ttcatgtttt acaatgatgt taagaagcag caacaaatgg cagcacttac
                                                                 2700
                                                                 2760
aaaagaaaag gaccaattat ctcagtctat tgttatgtat aaaagtttat ttgaagccag
ccaacagctt cttaatgaaa tgaaatgtca agttgaagaa gcaagattaa aagaggccca
                                                                 2820
2880
                                                                 2940
tattgaagca cttctgaaaa gaaagctatc agaacaagaa gaactgaaga aaaaacctag
                                                                 3000
aagatcgtgt actcttccaa actatactaa aggcagtgga gatgttttgg gaaagattgc
acatetagea caaattgaag atgatagage tgegatggtt atttettgge atetggeaag
                                                                 3060
                                                                 3120
tgacatggac tgtgtagtca ccctaaccac tgacgctgca cgtcgtatct atgatgaaac
                                                                 3180
ccaaggtcgt cagcaggtgt tgccccttga ttctatttac aagaagactc ttccagattg
                                                                 3240
gaaaagatet etaceteatt teegaaatgg aaaattgtat tttaaaecea ttggagatee
                                                                 3300
agtctttgct cgagacttgt taacatttcc agataatgta gaacattgtg aaacaggttg
                                                                 3360
ttaaaattac acactgtcct acactgctga ccagagatgg agatcgaatt cgaagtaatg
                                                                 3420
gaaagttigg gggccticag aataaagctc ciccaatgga taaacticgg ggaatggtat
                                                                 3480
ttggagetee agticeaaaa cagtgtetga tettagggga acaaatagat ettetteage
                                                                 3540
agtategite igeigtgige aaactagaca gigigaataa ggatettaac agteaattag
                                                                 3600
agtaccticg cacteeggat atgaggaaga aaaagcaaga actigatgaa catgagaaaa
                                                                 3660
atctcaaact aatagaggaa aaactaggta tgactcccat acgtaagtgt aatgactcat
                                                                 3720
tgcgtcattc accaaaggtt gagacgacag attgtccagt tcctcctaaa agaatgagac
                                                                 3780
gagaagctac aagacaaaat aggattataa ccaaaacaga tgtatgagag gtgacagaga
                                                                 3840
gaagaggcca ttggtctcag taagaatgcc ctgctttctg catctctgtt tcagaagacc
aagagggtga cttaccagac tgagtatttc tggggacaat acaagtacct gggcatgaat
                                                                 3900
                                                                 3960
ttccatttcg attcagatgg gactggaaac aaccattcaa ttttatgaat cttactggac
attatggatt tactggaatt attccagaca ttatgccctt tggttgtcac taccttgcaa
                                                                 4020
                                                                 4080
atgigiaaga ggaaaatgig claatgiggc agigactgia aaaciggcac atggcattia
ltaatcciga agaaaagtac aigtactati titcagtata aatataatga acatgicaga
                                                                 4140
                                                                 4143
act
```

<211> 3301

<212> DNA

<213> Homo sapiens

<400> 2029

atataggagg tggtttgctt ttgttgggca gtttatcacc ttcatgacca ccacaacacc 60 tttgctgttg gctccacacc cacagtcagt tttaacagga gtttcagtga atcagttagt 120

tgtaaccaaa	ggagttgccg	gccttcagtt	tattggattc	ggtgctgtgt	gtctgcctat	180
tcctcttgat	ggggaaactg	gagcagttcc	ctacagtcca	gccatttcag	gtgcccaatt	240
atgtctcctc	tacctgtgat	gttcagagat	gagaagagcc	acttttactt	tttcactgta	300
aatttttatt	taatgtcagc	cttgcttgcc	gaactataaa	ctctgtgagg	aggtctgtag	360
tgctcaccat	tgtttcttta	gagctgaata	cgtagcctga	cacacagtag	gctttcaata	420
aaaatttaat	ttaccagaag	tggaaaatga	gttttatgaa	gaaaatttca	gaaaactgag	480
ttcatttttc	aacacaagag	atgaccaagg	ggtaatatgt	tccttcaggt	tcatgaacag	540
cctgcatgaa	tatgccaagt	agttgttttg	taactgtgga	agattggcta	agaggagatg	600
gatggaaagt	aaagtcagaa	agaccttatt	gatttaggcc	agtgggagaa	gtgttggagt	660
atctgctctg	gagaaaatgc	tcttttccgg	ctagttttgt	taattatgtt	tctgaaaagg	720
ggggctagat	tggatggtct	ttaccaggtt	tcttcccctt	ctgattcagg	gacttcagga	780
ggtttgtggt	aacctgagaa	agtagcctga	ggtattatgg	tgctggagtt	ctccataggg	840
tgcttagcag	accaccttta	tctccccata	cattgcgttt	ttccatatgt	gagctgagaa	900
taagctggtt	gcctttcagt	gatctgaaat	tatagatgca	tttcttggaa	gctttatttt	960
ttttaatggc	taaaattgag	tagtatcgct	attgctgtct	gtagactacc	acttgctatt	1020
cctgtttaga	gtttactggg	cttggtaagt	tggaagggta	acaggagcac	gtttgtgatt	1080
ttttttttt	tttttttgag	acggagtctc	gctctgtcgc	ccaggccgga	ctgcggactg	1140
cagtggcgca	atctcggctc	actgcaagct	ccgcttcccg	ggttcacgcc	attctcctgc	1200
ctcagcetee	caagtagctg	ggactacagg	cgcccgccac	cgcgcccggc	taatttttg	1260
tatttttagt	agagacgggg	tttcaccttg	ttagccagga	tggtctcgat	ctcctgacct	1320
catgatecae	ccgcctcggc	ctcccaaagt	gctgggatta	caggcgtgag	ccaccgcgcc	1380
cggcccacgt	ttgtgattta	aacaacaaca	acaacaacaa	caaccagtta	acgtaattga	1440
cagcagagaa	gttccaggca	gaacagtggc	tctttcgttt	ttcttctaca	catggctttt	1500
tgccatcagc	atcagtgaag	acttgcggaa	ggagctaatg	ctgcttattt	gcagttgttg	1560
aacctgtttg	cctatgggac	atacccagat	tacatagcca	acaaggagag	cctgccagaa	1620
ctgagcacag	ctcagcagaa	caagctgaag	catcttacca	tcgtgagctt	ggcatcaaga	1680
atgaagtgta	tecectacte	cgtgttgctg	aaagacctgg	agatgcggaa	tctccgggaa	1740
ctagaagacc	ttatcattga	ggctgtctac	actgacatca	tccagggcaa	gctggaccag	1800
cgaaaccagc	tgctggaagt	ggatttctgc	attggccgtg	acateegaaa	gaaggatatc	1860
aataatatig	tcaagaccct	gcatgaatgg	tgtgatggct	gtgaagcagt	tctactgggc	1920
atcgagcagc	aagttctgag	agccaaccag	tacaaagaga	accacaaccg	aactcagcag	1980
caggtagaag	cagagattgc	ttgttttcag	agggaaaaaac	gtgatgtccc	cctcctgaat	2040
cttataacaa	cagetttett	ctggttacca	acatcaagaa	gacactcaaa	gccaccgcat	2100
cctcctcgge	tcaggagatg	gagcagcagc	tggctgaacg	ggagtgtccc	ectcacgetg	2160
agcagaggca	gcccaccaag	aagatgtcca	aagtgaaagg	tctggtctcc	agccgccact	2220
agggccggct	ggggcagctg	gcactcacca	ggcctgggtc	aggtggggag	gggacaccaa	2280

```
2340
gggcccattt cctcccctct ctacctgcag tgagttccag acctgcccgt cccctcacca
                                                                   2400
gegeeteece accetgttgg tactgtteea gaaaaactgt tacteecect cacceactee
                                                                   2460
ctccttcccc agttgttccc ttcagactca ggggctccac caatgccatc ccaaaacagg
                                                                   2520
gtcagacact gcccagcttc cctccaggag gttcttgtct ctgtgtaagg gcttgtctcc
ctcccagttt ttcttttgct ccacgtcatt ttgtcaggct ggttataagc cggaggcagc
                                                                   2580
                                                                   2640
ttlaaccage eeccagggat gattgtgaag gaggeeeete eeettgtgag gagggggae
tectetecag eccetggtae caeagteete aegatggtge agtgatttet ageeaggegt
                                                                   2700
                                                                   2760
caagatgege tgettteeet eteetgeete ateeettgtt ggeageteea gtteaggeeg
                                                                   2820
tggagggacg tgatgctggg ctgtgtttac taaacccacg ggttttcagc ctcttaagcc
cagctccgat ctccaattag ttgagagcgc tgggttgact aacctctggt atctgagcac
                                                                   2880
                                                                   2940
agacagaggg tgctgtgggt ctgctgggtg gcagaaatgg ttccttccgg cttggcgttc
telectggee actetteetg etgeetetga etacteagee ttgttttegg tgtgtaggee
                                                                    3000
                                                                   3060
ccagctgccc actggaactg ccggctaatg cttgctctcc caagatettt aactcctcct
                                                                   3120
ggctgcacct gggtagggat ggtggcatcg atgcccctct gtctgctgaa ggacctgttg
                                                                   3180
ctgcttctgt cttttcaccc ctccttggct gatgacccag agccctctga tgatggcatt
ctcctggcaa gagaaaaaga cttaactaga cttctgaact tgaacagttt caggttatat
                                                                   3240
tttaattttt ttttttttg tacaggttct gattctaata catttcaaca tgcttttgtc
                                                                   3300
c
                                                                    3301
```

<211> 3484

<212> DNA

<213> Homo sapiens

<400> 2030

60 attgcaaage cacagggaac ataccaggtg ttacctccaa acacaggete tetttatait 120 gacaaggatt gttcagctgt gtactgccat gagtcaagca gtaatatata ctatcctttt caaaagegtg ageagetgeg agetggeagg tacateatga ggeatactte agaggttate 180 tgtgaggtte tggateetga gggaaacaet ttteaggtea tggetgatgg tageatatea 240 300 actatattac ctgaaaaaaa attggaagat gatttaaatg agaaaactga gggctatgat agtotgtoot ctatgoacci tgaaaagaat catcagcaaa totatggtga acatgtoocc 360 420 aggittittg tiatgiatgc igatggatca ggaatggaac ticticgaga cagigacata gaagaatate tatettigge atalaaagaa teaaataetg tigiteteea agageeagig 480 caggaacagc caggcaccct aaccatcaca gtccttcgcc ctttccatga agcatcacca 540 600 tggcaagtaa aaaaggaaga tacaatigte eeteetaate teeggteaag gteatgggaa

```
acatttccct cagttgagaa aaaaactcca ggacctccgt ttggtactca gatttggaaa
                                                                     660
                                                                     720
ggcctttgca ttgagtccaa acagctagtg agtgccccgg gtgccatact caagagcccc
                                                                     780
agtgtgctac agatgcgcca attcattcag catgaggtca taaagaatga ggtgaaactg
                                                                     840
aggctgcagg tttcccttaa ggaltacata aactalattc taaagaaaga agatgagctg
caggaaatga tggttaaaga ttccagaact gaggaggaga gaggcaatgc tgctgatctc
                                                                     900
ctcaagctgg ttatgtcttt ccctaaaatg gaggaaacta caaaaagtca tgttactgaa
                                                                     960
gttgcagete acctaactga tttattcaag cagtettigg etacgeetee aaaatgeeca
                                                                    1020
                                                                    1080
ccagacacat ttggtaaaga tttctttgaa aagacatgga gacacacagc atcctcaaaa
cgctggaaag aaaagataga caaaacgagg aaggaaattg agacaacaca gaattaccta
                                                                    1140
atggatatta agaaccgcat aataccaccc ttttttaaat ctgaattgaa ccagttatat
                                                                    1200
                                                                    1260
cagtotoagt ataatoacot ggacagtott tocaaaaaac tgoottottt tacaaagaaa
aatgaagatg caaacgaaac agctgttcaa gatacatctg atcttaatct agatttcaag
                                                                    1320
                                                                    1380
ccacataagg titcagaaca gaaatccica agigigccia gicticcaaa accagagati
telgeagata agaaggatti caetgeteag aaccaaactg aaaatttaac aaaateteet
                                                                    1440
                                                                    1500
gaagaagcag aatettatga geeegtgaaa atteeaacee agteettget geaggatgtt
gegggacaaa caagaaaaga aaaagtgaag ttgeeteatt atttgetgag ttecaageet
                                                                    1560
aagtotcaac otottgoaaa ggtgoaagat totgttggag gaaaagtgaa cacatootot
                                                                    1620
gtigcatetg etgecattaa taatgeaaag teateeettt tigggiteea teeteteea
                                                                    1680
tcatcagtca agtttggagt gcttaaggaa ggacatacct atgccacagt tgtaaagctc
                                                                    1740
aagaatgttg gagtggactt ctgcaggttt aaagtaaagc agcccccacc cagcacagga
                                                                    1800
                                                                    1860
ctgaaagtga cttacaaacc tggacctgtg gcagctggta tgcagacaga actgaatata
gagttattig ccacagetgi tggagaggai ggggccaagg gaicagcaca caicteteac
                                                                    1920
                                                                    1980
aatategaga tiatgacaga geatgaggit etgiteetae etgitggaage aaatateett
taaagticaa citgagtaat catatatagi gcagaaalta cacgagigag gaaaacaigg
                                                                    2040
aagtcaaaat gcatctetac titaltaatt cialciicaa aatcagagit aaatttatta
                                                                    2100
                                                                    2160
agacaaagag catcitcati catciilgaa agcacciagc caaatciaaa aaaataccig
                                                                    2220
acacatagta tatgtgcagt aacttcagat tgaataaatg taaatgttat tggctatcta
                                                                    2280
eggaatatea gacagaataa taaaacagca agtatetate acaaaaaaat tataatttta
tggaaggata ggaaatacct taltattata aaggitgggi attcactgaa ttatgcatgc
                                                                    2340
attectectt ateagtgtet teagecaaac agatattaga tagatateaa gaacetatta
                                                                    2400
ectecaaggt actgtataaa atagtttate atatataaaa atggataatt ggactetgte
                                                                    2460
ttaaaaggta ttatataatt tgtagcagaa ataaagtett cacattttat ttetattttg
                                                                    2520
                                                                    2580
tactitetee agiggeatga attgigiget geligigila eagileteia titatitgat
ttttgagctg gatcttatag aatgigaaaa ctigaitgac gggaacttta agtaaaaata
                                                                    2640
                                                                    2700
atgaacaaaa ccatggcaac aggaaagctc caggtgtttg ggatgattgg cagggagttc
aacttgccaa aagcttgagt attaggaata tagtgggaaa gtaggttgga gtcaagttat
                                                                    2760
```

gaaagatctt aaatccttgg	cttgaatttt	attatttaag	cagcagtgaa	ccactgcaga	2820
ttcctgaccc tgtgggtgac	atgatcagca	tatctttatt	aagatgaatc	cagggttatt	2880
gtgcaggaca tgtcaaaggg	gaacaactgg	atgtgtaaaa	gtaccattag	aagtctacct	2940
gaatgggcca tgtgtgagga	caagaactgg	gagtggggga	acagtcaaca	taaaagaggg	3000
acatgaatga aagacatggt	gggggaagga	aactgcaaaa	tctgaggtag	aagccattga	3060
tggatggaag aaagaggaca	tcgagttcaa	cttcaaagtt	ttgggctgag	gtaatgaatc	3120
atgtatatgt aatattagat	ctcaactgag	aagtcagaat	tggagatata	ataattttaa	3180
gcatcgttta cacagaggtg	atggctgaat	gtatgggcaa	ggaacagaaa	tctggagtcg	3240
gtttagggag caggaggaag	aagagccagt	ggagacaaaa	gcagcaatta	gaaaatggtg	3300
aaatacttca gaagccttag	gaaaaatttc	aaggaaaaga	cggacacaat	tgacggatgc	3360
tattgagatg tcaaagaaaa	ttcagattta	aagtgttaaa	tttggttggg	ataaaaacta	3420
aattgcaaaa ggtaaagaat	gactgtatia	agaaagcaga	aacattagtt	atggatattc	3480
tttc					3484

<211> 3635

<212> DNA

<213> Homo sapiens

<400> 2031

ctititagag aatcttattc ccaaatattt gactccigag gicalicagg aagaalicag 60 120 teacatgett atatgeagag eaggagegee agettetega eatgetgtga aggtggteea 180 gaagtgtaaa atacaaaaag tgagattcca gggaaagtgc ccaccaagat caaggatatc 240 tgtgccaatt aaaaggaatg ctatattgca tagaaatgaa tggagaccac cagctggagc 300 ccagaaggcc agatctataa aaatgataga aagacccaaa attgctgctg tctgtggaca 360 ttatgattat tattatgete aacttgatat getgaggagg agageeeaca aaccaagtta 420 tcaccctatt cctcaagaaa atactggagt tgaggattac ggtcaggaaa cgaggcatgg 480 tecatececa agteaatgge etgetgagta eetteagaga aaatttgaag eteaacaata taagttgaaa gtggagaagc aattgggtot togtocatot totgoogago caaattacaa 540 600 cegagacaag agctaagaag taatggagaa gagcetagat teeaggaget gecatttagg aaaaacgaaa tgaaggaaca ggaatattgg aagcagttag aggaaatacg ccaacagtac 660 720 cacaatgaca tgaaagaaat tagaaagaag atggggagag aaccagagga gaactcaaaa ataagtcata aaacctattt ggtgaagaag agtaacctgc cigiccatca agaigcatci 780 840 gagggagaag cacctgtgca gaaggaattt cgctcttgtt gcccaggctg gagtgcagtg 900 gegegatett ggeteacege aaceteegee teecaggite aagegaliet ceigecteag

cctcctgagt	agctggaatt	ataggcgcct	gccaccgcgc	ccagctaatt	tttgtatttt	960
agtggagaca	gggtctcacc	atgttggcca	ggctggtctt	gaactcctga	cctcgggtga	1020
tccacctgcc	tcagcctccc	aaagtgctgg	gattataggc	atgagecace	ccgcctgagc	1080
gaattattat	tatctttata	attagagtaa	ttctctgtgt	tttaaattat	atttattatt	1140
agagcttggt	ccagagtcaa	ctagaaatgg	aaaatcctca	aggtattata	aacttgtcat	1200
ttaaaggtgc	cagtaggatc	acagtcacat	tccataaaaa	cacggctcag	atgitacaga	1260
catgtttttc	tctcacattt	tttaacctgg	ttagagtaaa	tccagtgcct	taaagttttt	1320
aataagtcag	gtaattaaaa	ataaaccact	ggaagcctca	aaaagtttgt	atcaggaatt	1380
gggtgaataa	aatcttgtat	attttatgca	agaggagtaa	ctttgaaaga	aaacacacca	1440
aaatgccaat	ggtggtaatt	ggtggtatct	ggattggtgt	gagtaggaat	gattattgtc	1500
tctctacttt	ttagattttt	tataagaagg	ttacagaact	tttactacaa	atatgtataa	1560
taaagtatcc	gttccttagt	tctgtcagca	ctctaatcaa	tatcttcaaa	caaaaaagcc	1620
atctgaaaga	cagaaatggt	ggcacgagac	tatagttcca	gctatttagg	aggccgagga	1680
tcccttgagc	tcaggagttt	gagaccagcc	ttggtaatat	agtgagaccc	catctctaaa	1740
aaaaaagaaa	aggcatctga	tatttcctga	aggctcctcc	agagcaatcc	agcagcagat	1800
acctttgcaa	acttttgtaa	aggaaataat	tatcacttaa	tttgtctaat	ttttggattt	1860
aggttttaat	tatcttttt	gaagggaata	tgcagctata	taataagaca	ctttaaaaaaa	1920
gtctctactt	gtagagttat	ctttccaaaa	tactgatttg	aacattattt	ctctacacga	1980
caatcaatgg	cgactgccat	ttctcttagc	atggcatgct	agacttttgt	gagttgttcc	2040
taacagaatg	ttccagcctc	attgctcaca	tttcccccaa	acatacccaa	agctctaaat	2100
gtctcagatt	acctttttt	tttttaaatg	acatatttt	tatttcttta	agtgatttt	2160
ttcactgtgg	taaaatacat	ataacatcgc	ctttaccacc	ctaaccattt	ttttttttt	2220
tttttttaat	tgatcattct	tgggtgtttc	tcgcagaggg	gtatttggca	gggtcatagg	2280
acaacagtgg	agggaaggtc	agcagacaaa	caagtgaaca	aaggtctctg	gttttcctag	2340
gcagaggacc	ctgcggcctt	ccgcagtgtt	tgtgtccctg	ggtacttgag	attagggagt	2400
ggtgatgact	cttaacgagc	atgctgcctt	caagcatctg	tttaacaaag	cacatcttgc	2460
accgccctta	atccatttaa	ccctgagtgg	acacagcaca	tgtttcagag	ggcacagggt	2520
tgggggtaag	gtcacagatc	aacaggatca	caaggcagaa	gaattittet	tactatagaa	2580
caaaatgaaa	agtctcccat	gtctacctct	ttctacacag	acaeggeaac	cateegattt	2640
ctcaatcttt	tccccgcctt	tccctcttt	ctattccaca	aaaccgccat	tgtcatcatg	2700
gcccgttctc	aatgagctgt	tgggtacacc	tcccagacgg	ggtggtggcc	gggcagaggg	2760
gcttctcact	tcccagtagg	ggcggccggg	cagaggcgcc	cctcacctcc	cggacgaggc	2820
ggctggccgg	gcggggggct	gaccccccc	cacctccctc	ccggatgggg	cggctggccg	2880
ggcggggggc	tgacccccc	ccacctccct	cccggacggg	gcggctggcc	tggcgggggc	2940
tgacccccac	ctccctcctg	gacggggtgg	ctgccgggcg	gagacgetee	tcacctccca	3000

gacggggtgg	ctgccgggcg	gataggctcc	tcacttctca	gaccgggcgg	ctgccgggcg	3060
gaggggctcc	tcacttctta	gacggggcgg	ttgccaggcg	gagggtctcc	tcgcttctca	3120
gatggggcgg	ccgggcagag	acgctcctca	cctcccagac	agggtcgcgg	ccgggtagag	3180
gcgctcctca	catcccagac	ggggcggcgg	ggcaaaggcg	ctcccacat	ctcagacgat	3240
gggcggccgg	gcagagacgc	tcctcacttc	ctagatggga	tggcggcggg	gcagagacgc	3300
tcctcacttt	ccagactggg	cagccaggca	gaggggctcc	tcacgtccca	gacgatgggc	3360
ggccgggcag	agacgctcct	cacttcccag	acggggtggc	ggccgggcag	aggctgcaat	3420
ctcggcactt	tgggaggcca	aggcaggcgg	gtgggaggtg	gaggttgtag	ccagccgaga	3480
tcgcgccact	gcgctccagc	ctgggcacca	ttgagcactg	agtgaaccag	actccgtctg	3540
caatcccggc	acctcgggag	gctgaggctg	gcggatcact	cgctgttagg	agctggagac	3600
cagcccggcc	aacacagcga	aaccccgtct	ccacc			3635

<211> 4050

<212> DNA

<213> Homo sapiens

```
60
aaatgttatt agttgctatg titgggtigi ggggtgatag gtgcttictg tilacticit
                                                                     120
tgtgctttct tctattttct gcaatgaatt tctgttttat cattagaaat aacagtaggt
                                                                     180
attitaaatt acacaatgaa ataaacaacc tagggcacac taaattigtc atggattetg
agetecaagg aacaggteag cettaceagg eccageetee eteeeetgea getgtgggge
                                                                     240
                                                                     300
ataggattet cagcaagtgg gtacagatgg aaataccagt gcagtggete tattetgatg
                                                                     360
tggactgaag aggccagatg ggaaacatcc tattccaacc tggactcttc ctgcaaggag
                                                                     420
gatgccaacc aactggaggc ccctggagaa aggacaccag gatggaggga gtgacactcg
                                                                     480
aggtcatggg caggtttatt cittaaagig cagtcatggg ggaggtggga agacacagic
ttgatettea aateteaaga giicialeet gggeagagae ageaaciitg eiitteatei
                                                                     540
                                                                     600
ccacaaagga cagacctagg acaaatgtga gacagattgg agctcaggat gatagcaaat
                                                                     660
cagtgcagtc cccaggggga ggttgtatgg agacaaatta tatatttgtt titcaaacct
ggaaagagac aggagatgaa cagagtgttt tctttattta tttatgccct acatcttccc
                                                                     720
                                                                     780
ccaaaggatt ttaaattgtt tacacggaat agtatgtgga tcataatgtt aatggaattt
                                                                    840
aaattggaaa teagggeeaa agaaaggaga atgaageeaa tgttettetg tatgagetge
                                                                    900
taacgggctt gaatgtgett aattitgaac ctgagcticc tgtcatgctg cgtlagaaag
                                                                    960
aagaattgat ttgtgtatte atteaacaat atttatteaa gtatttatag ageacataet
atgtgccaaa cattgttcta gatatagagt aaagtgacca aacacaacgc accatagcac
                                                                    1020
```

ctcctctccg	ggagggaata	ttctagtgag	aaaagacaaa	taatacttga	aactgttgac	1080
aaagagagtc	aaactctgta	aaatacttga	agagatttat	tctgagccaa	atatgagtga	1140
acaatggcct	gtaatacagc	cctcaggaga	tcctgaaaac	atgtacccaa	ggtggttggg	1200
ccacaacttg	gttttataca	ttttagggag	atgtaaggca	tcagtcaata	catgtaagtt	1260
gtatttggtt	tggtctggaa	aggtgggaca	actggaagca	ggggctttca	ggtcataggc	1320
agattcaaag	attttctgat	tgacagttgg	ttgaaagagt	taagttattg	tctaaagaaa	1380
ggaatgtctg	ggttaagata	aggggttgtg	cagactaagg	tcttatcata	gagatgaagc	1440
ctcccggttg	taggcttcag	aggataggct	gtaaatgttt	ctatcagact	taaagagtct	1500
gttctaacag	taattccaaa	aaggaggagg	gtataatgaa	gtaggtttgc	cgccccttc	1560
ccatcatggc	ctgaactagt	ttttcaggtt	aactttggaa	tgcccctgac	tgagaggaga	1620
ggtccattca	gatggctggg	ggcttagaat	tttatttttc	atttatgaaa	cacaaaaaga	1680
agccaagaaa	tgaatgagct	tggaaaatat	tagacagtaa	taggcactga	gtgaagatat	1740
cggggggacc	aatgtcacca	ggaggtgaca	tttaagctga	ggtctgagtg	aaaagaaaca	1800
gactttgagt	gacaataatt	ttataacaaa	cactggaggc	agtttttcca	gggactgttt	1860
ttggaaccag	cctccagagc	aaagaatctg	ccttttaggc	gcagttcagc	aaaggggtgt	1920
tgtaaggtca	gggcccgtgg	gccctgcttg	tgcaggcttc	tggtggtccc	acgaaattcc	1980
agaagaaaaa	actggagtcc	tagctgaaca	atgtgtgcct	cagcaactgt	cttcctggag	2040
ttttcctttt	ctcagctggg	cttttgatag	gagtccagta	gcagatacct	ggagagtttg	2100
ttgcacgaag	aatggctgcc	caccattgtc	aactttgtct	ctatccttct	ctgaatgaag	2160
agaactagag	cacatctaat	gttgtcccta	ctcaactgac	caccttgcat	tggaggaact	2220
tgttttgagt	tacataatta	ggctaagaga	aacaaaccta	gaaacctggg	ttcctcattt	2280
gttgcaacat	tcctcaaggt	tctctctggc	agaagccata	cgataaaata	tctttaaatt	2340
gggcaacctg	gcttttcatc	ccagccagct	gtgtgatttt	gggttggtga	ctaatttgtg	2400
ttttccacat	taatacagtg	agaaggatta	tttttgttct	gcctatatcc	tagggacttc	2460
atatggaaga	agtaaagtga	cagctgggaa	agggacttta	ggtgtcaacg	gcagtatgag	2520
aatacaggat	ttttgtcaat	ctgctgtgtt	tccccaggtt	aggaaaaaccc	tggatgccac	2580
catgcagaca	ttacaggaca	tgctgactgt	ggaggacttt	gatgtctccg	atgccttcca	2640
acacagtcga	tcgacagagt	ccgtcaagtc	ggctgcctct	gagacctaca	tgagcaagat	2700
caacattgcc	aagaggagag	ccaaccagca	ggaaacagaa	atgttttatt	ttacagtaag	2760
tggcatcctg	ggcccagaac	cacactgtcg	gccaagccac	tggcagtgac	ttttcaggag	2820
caacccaagc	tactgagaac	cagagaaacc	acatgggtca	attggatcta	agactccatc	2880
accatgcttt	taaaattaag	ttgcctggct	tggttttctg	aaatgcagaa	agtggattcc	2940
caatgggtag	cattggcatt	gatcttgggt	gatgattatt	gaaattttct	tgctctagaa	3000
aaaaaccaga	gacagtttta	ttcagtgggg	tgataagaaa	atggctgaca	gagtcaggta	3060
caagtcccaa	ggaacaacct	tgaaattatg	tatatagatt	atcatgttga	attgtcttaa	3120
atttaggtgt	gagctttgga	aaaaatgccc	tcaaaaatcc	aagcaaattg	ctcttgagtt	3180

gctagccctt	catgtaaaat	cccatgttaa	ttatctttca	tttggacagg	gactgggagg	3240
agaaaggaga	cggggactgg	ggctttagtt	caacatgtgt	ttactgaaca	tagaatatta	3300
ggtttgtaag	ggaccttaaa	tcttccatgg	gatgcttgag	tcagttcagc	cacatccctg	3360
accaggggcc	atcctgactc	tgcctgattg	ctcccaagat	taaaatctcc	ctttttcccc	3420
agacagccct	tttgtttgca	ttgtgcctgg	cttttggtat	taccatgttt	tcctttatct	3480
ttgctatctt	caaaaaccta	cctcactagg	atttcttggt	tctgttctct	ggggccaccc	3540
agagtggagg	ctaattctac	atggcagtgt	ttcacatggt	tgcagggagc	tgggatttca	3600
tttctctagg	ctaaatgtat	ttgattcttt	cagtcttgcc	ttataagctt	ttgttttgag	3660
tagcctcagt	atcctagtga	ctctctctg	gacatgttcc	atgtgctgat	gcctcttcta	3720
aggtgagact	ccgagcagtg	gtatgcccaa	cacagaactg	agcaaaattg	gccgggcacc	3780
gtggcttacg	cctgtaatcc	gagcactttg	ggaggccaag	gcaggcagat	tgcctgagct	3840
caggagttct	agaccagcct	gggcaacacg	gtgaaaaccc	gtctctacta	aaatacaaaa	3900
aattagctgg	gtgtggtggc	gtgcacctgt	agtcccagct	acttgggagg	ctgaggcagg	3960
agaattgctt	gaacctggga	agtggaggtt	gcagtgagcc	aagatcgtgc	cactgcctcc	4020
agcctgggtg	acagagcgag	acttcatctc				4050
	agaaaggaga ggtttgtaag accaggggcc agacagccct ttgctatctt agagtggagg tttctctagg tagcctcagt aggtgagact gtggcttacg caggagttct aattagctgg agaattgctt	agaaaggaga cggggactgg ggtttgtaag ggaccttaaa accaggggcc atcctgactc agacagccct tttgtttgca ttgctatctt caaaaaaccta agagtggagg ctaattctac tttctctagg ctaaatgtat tagcctcagt atcctagtga aggtgagact ccgagcagtg gtggcttacg cctgtaatcc caggagttct agaccagcct aattagctgg gtgtggtggc agaattgctt gaacctggga	agaaaggaga cggggactgg ggctttagtt ggtttgtaag ggaccttaaa tcttccatgg accaggggcc atcctgactc tgcctgattg agacagccct tttgtttgca ttgtgcctgg ttgctatctt caaaaaaccta cctcactagg agagtggagg ctaattctac atggcagtgt tttctctagg ctaaatgtat ttgattctt tagcctcagt atcctagtga ctctctcctg aggtgagact ccgagcagtg gtatgcccaa gtggcttacg cctgtaatcc gagcactttg caggagttct agaccagcct gggcaacacg aattagctgg gtgtggtggc gtgcacctgt	agaaaggaga cggggactgg ggctttagtt caacatgtgt ggtttgtaag ggaccttaaa tcttccatgg gatgcttgag accaggggcc atcctgactc tgcctgattg ctcccaagat agacagccct tttgtttgca ttgtgcctgg cttttggtat ttgctatctt caaaaaccta cctcactagg atttcttggt agagtggagg ctaattctac atggcagtgt ttcacatggt tttctctagg ctaaatgtat ttgattcttt cagtcttgcc tagcctcagt atcctagtga ctctctcctg gacatgttcc aggtgagact ccgagcagtg gtatgcccaa cacagaactg gtggcttacg cctgtaatcc gagcactttg ggaggccaag caggagtct agaccagcct gggcaacacg gtgaaaaccc aattagctgg gtgtggtggc gtgcacctgt agcccagct agacttgcagattct gaacctggaattgct gaacctggaattgct gaacctggagattct gaacctggaagtggaggtt gcagtgagcc	agaaaggaga cggggactgg ggctttagtt caacatgtgt ttactgaaca ggtttgtaag ggaccttaaa tcttccatgg gatgcttgag tcagttcagc accaggggcc atcctgacte tgcctgattg ctcccaagat taaaatctcc agacagccct tttgtttgca ttgtgcctgg cttttggtat taccatgttt ttgctatctt caaaaaccta cctcactagg atttcttggt tctgttctct agagtggagg ctaattctac atggcagtgt ttcacatggt tgcagggagc tttctctagg ctaaatgtat ttgattcttt cagtcttgcc ttataagctt tagcctcagt atcctagtga ctctctcctg gacatgttcc atgtgctgat aggtgagact ccgagcagtg gtatgcccaa cacagaactg agcaaaattg gtggcttacg cctgtaatcc gagcactttg ggaggccaag gcaggcagat caggagttct agaccagct gggcaacacg gtgaaaaccc gtctctacta aattagctg gtgtggtgg gtgcacctgt agtcccagct acttgggagg agaattgctt gaacctggga agtggaggtt gcagtgagcc aagatcgtgc	getagecett catgtaaaat eecatgttaa ttatettea tttggacaag gaetggagggaaaagaaagaaa eggggactgg ggetttagtt eaacatgtgt ttaetgaaca tagaatatta ggtttgtaag ggacettaaa tetteeatgg gatgettgag teagtteage eacateeetg accaggggee ateetgaete tgeetgattg eteceaagat taaaatetee etttteeetg agacageeet tttgtttgea ttgtgeetgg ettttggtat taeeatgtt teetttatet ttgetatett eaaaaaceeta eeteactagg atttettggt tettgteet ggggeeaeee agagtggagg etaaateta atggeagtgt tteaeatggt tgeagggage tgggatteea tteeteteagg atteeteteagggtgagg etaaatgtat ttgatteett eageettgee ttataageet ttgttttgag tageeteegg ateeteetg gaeatgtee atggetgat geetetteta aggtgagaet eeggaagtg gtatgeeeaa eacagaaetg geaggeagat tgeetgget eaggagteet agaeeetgg gtgaaeaeeg gtgaaaaeee gtetetaeta aaataeaaaa aattagetgg gtgtggtge gtgeaeeetg agteeeage aggeetgee eacggagggaggaggaggaggagaattgeet gaaceetgggaggaggaggaggaggaggaggagaattgeet gaaceetgggaggaggaggaggaggaggaggaggagaattgeet gaaceetgggaggaggaggaggaggaggaggaggaggagaattgeet gaaceetgggaggaggaggaggaggaggaggaggaggaggaggagg

<211> 3663

<212> DNA

<213> Homo sapiens

```
gcgtgtggtt cttggagaaa gttggaggtg gtggtgattt cagtcgcctt ggccgccttg
                                                                     60
                                                                    120
agceggaget gageggagge aetgggeega geetgettee egggeettee taceatgeea
gggctgctcc ctgcctccgc caccttggca caccttcacc cgcgtaccgc ctcctcccg
                                                                    180
                                                                    240
tegetetgee titteeaaaa eteaetiggg eeeleegige geagggitet tittiggiit
ttetgtaaaa atcaaaacaa aaaacagaga ettttgagag gagcagatge cacctaaagt
                                                                    300
cccactgcat tccctgcaaa gcgctcaaat gtggaagcca gtcattggca tttttatttt
                                                                    360
                                                                    420
ttaltgattg attgatittt tcaccagigg ciltiigtaa ccicigigii cigcigigii
tettgtgttt agtettegag tgettegaet gaecatgate eeetgggeee eeteceteet
                                                                    480
                                                                    540
ggctgggaga agagacagga caatggacgg gtgtattacg tgaaccataa cactcgcacg
                                                                    600
acceagtggg aggateeeeg gaeeeagggg atgateeagg aaccagetet geeeecagga
tgggagatga aatacaccag cgagggggtg cgatactttg tggaccacaa tacccgcacc
                                                                    660
                                                                    720
accaccttta aggatecteg eeeggggttt gagtegggga egaageaagg tteeeetggt
gettatgace geagttiteg giggaagtat caccagitee gitteeteig ceatteaaat
                                                                    780
```

```
840
gecetaceta gecaegtgaa gateagegtt tecaggeaga egettttega agatteette
caacagatca tgaacatgaa accetatgac etgegeegee ggetetteat cateatgegt
                                                                     900
ggcgaggagg gcctggacta tgggggcatc gccagagagt ggtttttcct cctgtctcat
                                                                     960
gaggtgctca accctatgta ttglttattt gaatatgccg gaaagaacaa ttactgcctg
                                                                    1020
cagatcaacc ccgcctcctc catcaacccg gaccacctca cctactttcg ctttataggc
                                                                    1080
agattcateg ccatggcget gtaccatgga aagttcateg acacgggett caccetecet
                                                                    1140
ttctacaagc ggatgctcaa taagagacca accctgaaag acctggagtc cattgaccct
                                                                    1200
                                                                    1260
gagttetaca actecattgt etggateaaa gagaacaace tggaagaatg tggeetggag
                                                                    1320
ctgtacttca tccaggacat ggagatactg ggcaaggtga cgacccacga gctgaaggag
                                                                    1380
ggcggcgaga gcatccgggt cacagaggag aacaaggaag agtacatcat gctgctgact
                                                                    1440
gactggcgtt tcacccgagg cgtggaagag cagaccaaag ccttcctgga tggcttcaac
                                                                    1500
gaggtggccc cgctggagtg gctgcgctac tttgacgaga aagagctgga gctgatgctg
tgcggcatgc aggagataga cagagcgact ggcagaagag caccatctac cggcactaca
                                                                    1560
ccaagaacag caagcagatc cagtggttct ggcaggtggt gaaggagatg gacaacgaga
                                                                    1620
                                                                    1680
agaggateeg getgetgeag tttgteaceg gtacetgeeg cetgeeegte gggggatttg
                                                                    1740
cegaacteat eggtageaac ggaccacaga agttttgeat tgacaaagtt ggcaaggaaa
cctggctgcc cagaagccac acctgcttca accgtctgga tcttccaccc tacaagagct
                                                                    1800
acgaacagct gagagagaag ctgctgtatg ccattgagga gaccgagggc tttggacagg
                                                                    1860
                                                                    1920
agtaaccgag gccgcccctc ccacgccccc cagcgcacat gtagtcctga gtcctccctg
                                                                    1980
cctgagaggc cactggcccc gcagcccttg ggaggccccc gtggatgtgg ccctgtgtgg
                                                                    2040
gaccacactg tcatctcgct gctggcagaa aagcctgatc ccaggaggcc ctgcagttcc
                                                                    2100
cccgacccgc ggatggcagt ctggaataaa gccccctagt tgcctttggc cccacctttg
caaagttcca gagggctgac cctctctgca aaactctccc ctgtcctcta gaccccaccc
                                                                    2160
                                                                    2220
tgggtgtatg tgagtgtgca agggaaggtg ttgcatcccc aggggctgcc gcagaggccg
                                                                    2280
gagacetect ggactagtic ggcgaggaga ctggccactg ggggtggctg ttcgggactg
                                                                    2340
agagcgccaa gggtctttgc cagcaaagga ggttctgcct gtaattgagc ctctctgatg
                                                                    2400
atggagatga agtgaaggte tgagggageg ggeeetgggg egaggeeate tetgeetgee
tecetageag gegeeagegg tggaggetga gtegeaggae acatgeegge cagttaatte
                                                                    2460
attoteagec aatgaaggtt tgtetaaget geetgggtat eeaegggaca aaaacagcaa
                                                                    2520
                                                                    2580
acteceteag actitigteea tigtataaact tigaagtigtit gigtigtagig gitigeaggit
                                                                    2640
ttttgttacg ctgctgtcac tttctgtcca ggagctggca ccccaggtgt tctgagacct
                                                                    2700
tgagggaccc agacctttgg gtccaagagt ttcccaaaca gccacgcctc tcaggaaccc
                                                                    2760
acctggcggt tccgtgagct caggcaggcc tgacccggcg gcacagcctg gcagggacct
                                                                    2820
cgtccccaag cctggcagaa tgagagggt tgaggtcccg agcgccactc ctagccttgc
                                                                    2880
cgccttcaat agagaagaaa tccctttgct agatagggtc ccccaggcag tccccagtgg
                                                                    2940
cgggacacag gggtccggct gtggagctcc cctgccagcc cctggagctc caggagggcc
```

tgttggtccc	ctgttcagaa	tggagtgcag	cccgccagcg	gaaagtgttc	attctgcata	3000
ggtgtgaggc	tttatctgca	cacaggacat	gaaaaccagc	agaaaggccc	tgagctgctg	3060
catagcccca	tctgatttct	gcagctcccg	ccagcctcca	acacggggac	tctgccgtaa	3120
ctggaatctt	cataggtcat	attgaaatct	tcaaggtgac	catgccccac	cggggtgctg	3180
gggcagtagt	catggcagac	tcccggcctg	ggcccccagg	attctaggac	ccccaggcag	3240
ccccttggac	tggtcccggg	tgccttccaa	gcacagtctc	catgctccca	gattctcgac	3300
cttccccgg	cccgggaggt	gcagcctgcg	tctgcctctg	tcgtgtgtgc	tgatttgagt	3360
ggcttagctt	gccacagcgc	agcctcttct	gtccctttca	gtcatttgct	gtacttccct	3420
gtggcacgtt	accatggaag	ccgctccagg	gtgggtcagg	gtgcaagctg	ctggtgaggt	3480
ttggaagcat	caggctcacg	ggtgttcatg	tgtgttcgtg	cgtgtgtgtg	cgtacgtgta	3540
tataactgaa	gtgtctgtac	ggaatgccct	ttgctagcca	tgggctggtc	accagattgt	3600
tttgtaatgc	ccgcccttg	cctcgatatt	gccagtttct	tgtgcaataa	acaatcagca	3660
gct						3663

<211> 3615

<212> DNA

<213> Homo sapiens

60	gctcgggggg	aggggttcgg	ggcagtggaa	gtgaggtgtt	cgggggcgag	aagatggcgg
120	ggcagtcggc	gcaagccccc	ttcacgctcc	gccgctcggc	cggtcctagc	cggggggacg
180	cgcacaccac	gcctggcagc	ggggggtcgt	gcacccgcgc	gtcaccaccg	aggaaccgcc
240	tatgggcctg	tgcctgtgca	ttggagaagc	cggtcgttcc	ggcgcgcgga	cggatgcgct
300	cgtggtgtgg	tcgcgagcct	ttctcggaca	ggaacccagc	aggtggagca	gtgatcaccg
360	tttcaaaaga	accaaggtat	gtctacgacc	ctacattagc	tgggcatctc	tgtatggccg
420	gggcctagat	aagaacttct	aaacaacagc	tgaaatttta	gattgatgga	aataattcca
480	agttttaaat	aagatgatca	agtaatgaca	atttgcaaat	actcaccaga	tgttcaaaat
540	tgtaagagct	aagcagatat	gaagatggaa	gctgtctccg	cagtgaaggt	tgccatttgg
600	tttggatgta	gacccacaga	aagcaaaaga	agtagcccag	tttgccagtt	gctcaggact
660	agtattgaag	atcctgattt	ggttgtcctg	tagttcaaat	ccagtttact	gatacgttag
720	gactgagatt	acatcagatt	cttccctggc	attaggcttt	tggacagcac	ttcggtcctg
780	ccttcgtcaa	ttttctctgc	tatgaggact	aaacatcagt	cttcccacct	gtctctttgc
840	atttgatttg	tggttgcata	tagtggtcat	tctgggaaag	gtgaacagcg	tatgcagcct
900	atgaaaccct	caaagcacct	ctgatgttta	ccaagtgact	aggaaaggaa	aggcttgtgg

```
960
gtacacacct agttcataat cetcataatt tatcaacaaa cacaaaaaag tgtcttactt
gagagtgagt gtgtgtgt gcgtgtgcac gtgcacacat gtgcacgttt gtatgtatgg
                                                                    1020
                                                                    1080
aaataaactt ataaatgggg acgtattgga gaaggaaata catagaccta caactttgag
caaatagcag tgatgtttta ggaactgaaa tgtcacactt aaagtcttca gcccagctac
                                                                    1140
                                                                    1200
ttccctattt ttgtggggag aagagggcct gattagaact gttctggttg tgtttggcgg
gaggggaata attitigite agicettett agigaceaaa ettiaattii taagaataat
                                                                    1260
                                                                    1320
atattgactt actgaactga agcattctga gttgaaagga gctccagagg agtggagttc
tgtgttgctc acatgttaaa atcttgctca ccttcagagc agagggaata cctatcttca
                                                                    1380
                                                                    1440
gatatccgtc cattiticatc tcttaattgt agtcaaaagt atgacttgag agtgttgctc
tggtattctg ggttctgaag tctggtattc tggtattctg ggttcaaaag tatgacttga
                                                                    1500
gagtgttgct ctggtattct gagagttgct ctgtattctg ggttctgaag attatttgaa
                                                                    1560
aaataactcc tactacattg aaatgcagac ttaaaaattt aaacattgga ttaggcagtc
                                                                    1620
aaaaaaacca agcaagcata aaaggtcaat aagttgtaat cttgatagta aaggtggaaa
                                                                    1680
                                                                    1740
acttattata aatggaaaga aagttttatt teetittittg titgatggge agtatgeeat
attataccca aagttettit aaaaaatatt teeatcaacc attittatti aaaataaaca
                                                                    1800
                                                                    1860
tttgagggaa gttaccaagg cagcittiit ccicaaaagi aaccigticc tcitiggaat
                                                                    1920
agcacatttt aggggcatgg ttaatacctg agatttttac tcagtaaatc ctgatggtta
                                                                    1980
ctgtgtgtaa aatatcttta agtaggattg aaggcctctg tgggggaata aaatattacc
aaagtctata aaaataaatt ttacatgttc tcttttatga cagagagcag cactggttct
                                                                    2040
gttattttta aaatgaataa ttgatttctt gataggtgtt taatatttct tccctcactg
                                                                    2100
ctgattetta gatagaaacc attetttata tttgatagac tgettteaga aaaccettat
                                                                    2160
                                                                    2220
caacaagtgt acaatactta tctaaaacta tacatttaga atggagcagt ttaatactag
atctcagaag tittgaaaaa tagcaaagaa gactggatti ggaaagcaig gictacaatt
                                                                    2280
                                                                    2340
ggttgttaaa ttctgaagct atgaagaata aatgtttcaa ctttggatta tgaaacccca
tttatgattt tttaaataca cttgaaataa aaatgattaa actaaatttt ggtccagtga
                                                                    2400
cattactttg cactgcataa tccattatac gttgtacgac ttttttttt gttttaattt
                                                                    2460
atlactgaga gttttgtgtg aagctacagc atatctaacc agagaatttc tgattcctta
                                                                    2520
                                                                    2580
tactgigatt atattatalt gaggcattig tagtgcagci gaagactgaa titatgcctt
                                                                    2640
ttglaaacat gataggtata aatgtettat aaacattetg gagtatgtat agetttaatg
                                                                    2700
2760
gtcaaataaa tlacctactg gaatatagcc caagccagta aaggtttaat atttgcattt
{\tt tcgtgctttt} \  \, {\tt altttctcct} \  \, {\tt tccattcata} \  \, {\tt agtatatact} \  \, {\tt tgaaagtaca} \  \, {\tt tctgtagcct}
                                                                    2820
                                                                    2880
atgalltigag tetettgaag tietaggaag aggeaaacta caaactacta ggattetgat
                                                                    2940
ticagatgia gicattccag aaccttctct ttatgagtic acctgetagt acaatctcca
caactigaat ggcattggti gitcigtaat teelgecaaa agcateacaa giigtacate
                                                                    3000
                                                                    3060
atcaaggete cettigeact eccaagaaga aciggtaatt ttaaacaaaa gtatgigtet
```

```
ttatttgtat tggaaaatac tgtctttaaa ttgtttcttg ttgacactcc ccacaatgga
                                                                    3120
aaaattaccg aattaaacct gttttatgga tggcagcttg gagcatagca agaagttgga
                                                                    3180
ggattigaat teeatteeea giteteatig igittigitt ettaaaacta taataategg
                                                                    3240
ttactgttat aaagtttaaa aggtggtctt aatgtgaata gcaaattctg gtatatcgtg
                                                                    3300
actaacgctt aagaatgcct gtctttgaga ggaaggtgtt ataatattaa tgaacagtgc
                                                                    3360
caaatacact gtgcatatct gcaatttaat ctttgaatgt atgttactgg attagctccc
                                                                    3420
tectectgtg tgatggtace atgeatagag teaateaaat cettgtgatg ttttgtatgg
                                                                    3480
actitgacaa tatgiaaata atgigtaaag ccagtittia tgattaagga atcaaattta
                                                                    3540
ttgaatttta ttattgaaag ttgaaactta acatgtatga acaaaaacca ataaaagaat
                                                                    3600
atactctttt cattg
                                                                    3615
```

<211> 3758

<212> DNA

<213> Homo sapiens

<400> 2035

ctgttgattg gccactgacc cgtgctgcag gcacacaaag gaagctgcac ccacagcagt 60 ctgttgtgga tggttgctga gctgcgcatt cggcattggg cttgctttgt ttcctgccag 120 geceageatt tietictace agateggeag getigiggge tieticetag gieceteece 180 tgcactciga ataggaaagc iggaagcigi gcittagaga agcittaaga cgccgaaaga 240 aaccagaaga gigagcgcca giigiatgig cgiggictcc atccgcaaag ccggagcigg 300 gcgcaacagt gttgacttgt aattgatcaa tttagatcgg gcgcaggccg ggggagggca 360 gtgcttttga tttaggctgg gaaaggcctc ctagtgacta tgttcaattt ggaggaattc 420 agatgetett tigitalaea agigaagetg tgiaalaeaa aigaggagii tiaetitiee 480 taaatcticc ccttatcalt caagtatiga ggagttttac ctttcctaaa tcttcccti 540 atcattccag tattatcagt gagatctggt tgtgatttat gtaaatggtg gctaaaaaat 600 tcaaactact gagggggaga attctcattt tacagcttca catgctgtgc tgaactaaat 660 aagtagegtg ggatgtigge ittgtgaeag gtetitigte attitieaga aageatitig 720 actigitgat gicaatiigg aacagcigaa aaaatacagg aaaataagat aaatacgiac 780 atgttgaggg tggggacaaa atgaaggttc tgaaccagct gccggcttac agtagccata 840 taagcaacag cagcaatgca ccaacciggi gagtaatagg ccigaticac iggagagata 900 ctagcacctt taatgagtca gatagatgca caatgggtgt gggagcagtt ggacttgtgg 960 gcacaaagtc tagcaagaag ctcagacttg caaacaactg taggacgtgc aaagcaagct 1020 ggcattggag cttgccgggc acagctgctc aggaataggc agctggtttt ccctttgatc 1080

```
1140
cctgagattc caaaggttac tttcctcttt gttcccttcc cagggtcaat tagagtagaa
actgcagatg cttttcagtt gagaattttc ctagaattct caaaaatgtg tatgctggct
                                                                     1200
                                                                    1260
taaaatetge catcaagaat tetgttacet tgetttaage etceagttee ttecagatgt
atggtggagg aggccagagg gcccttgttt tggggcttca gaggatggtt gttatctgga
                                                                     1320
                                                                    1380
tgagcactgt ggaaagactg agagagcaac tgagagaaag tgggcccctg aatgaaagtg
atttegeaaa ttttaggeag atgeeaceat cagaaactga tattttetga egtetttete
                                                                    1440
                                                                    1500
accttectet agageattea gtecagaaat gaccageetg tecaaagggg gaaattaetg
                                                                    1560
atattgatct gttccttaga gcagtgtttc agtctttttt ttttttttga gatggaatct
cattetgtca eccaggetgg agtgeagtgg caegateteg geteattgca acctecaect
                                                                    1620
                                                                    1680
tectgattea agtgattete etgeeteage eteceaagaa getggaatta eaggtgtgea
                                                                     1740
ccaccacac cggctaattt ttgaattttt tatagagatg gggtttcacc atgttgccag
                                                                    1800
getggtetea aacteetgae etcaagtgat ceteetgeet eggeeteeca aagegetagg
                                                                    1860
attacaggeg tgagccacca tggccggcct tcagcctttg tgatattaaa gcacagcaac
acattteeea ttacaeeeet gaacacacae acacagaaaa eecaaaagtt teacaaaatg
                                                                    1920
                                                                    1980
attettgete ttactactet cagtacacte tgtatttaaa aaaaaaaatg etggttgtgg
                                                                    2040
cttcctaagt ggtgcgtgca gttttcaaat caatgccctt ggcgataaag tgtgccctat
                                                                    2100
actgattate tetggacaaa gtetgaatgg ggettggete taatetetag teeteattgg
acattttaca tacctggcct ttgcctccac cctgatgtgg agtgatcatg ggggtgggaa
                                                                    2160
                                                                    2220
atatagctgg atccgaaagc tctgaagtgg ggatggaggt gtcacagctg aggctaggcc
                                                                    2280
cattetgeag ggeacteagt gtgtacagtt ggttttetat caggggteaa ceggegggg
                                                                    2340
gacitgagaa cagatetetg ggcacaaage agggeetttg ceetgggget tgetatgtgg
                                                                    2400
ctcagcctac acggetetet eccegteagt cetgtecaaa geecaggaaa etaatgtace
acccccgagg aagagagcct acctttccat ccaaggaagt gttttacctg tggtaagcac
                                                                    2460
                                                                    2520
gggggacaga attettgagg aaggagggtg etgegteeca gtggtggagg aaaagagagg
                                                                    2580
acciggtgia agcagccatg gcatggacci catccgaggi ggcacciggc tagggiccig
                                                                    2640
acctccaatc cttccccagt aaccatcact ttgagtaaac agtggctcca cccccggcat
ggttctttgc accaacattt ggggaatgcc taccaggggt cacacactga gctggatgct
                                                                    2700
gagtgtaggg tgtccacaac atcgtgccta aaaagtctct gtatggggta taagaaggtg
                                                                    2760
                                                                    2820
ctggggcaat acagatgaga tgagaagcat ctttcaggga atgggttgat cccaattcag
                                                                    2880
getteccaga gaaggatgte tgtagaette atattageaa gggaggaagg tagecaggee
                                                                    2940
acaggactge tggtgtaaag accagggcat atgaaatgge aagtgtgact gtgettteag
                                                                    3000
ccaataattt ggtattgtca aatgatggga ccaaacagct ggagaggcag atcctaaagg
                                                                    3060
gtcclgtggg ccaggclgga cttcatcttg tcactaacta alggagaggc tctgaaggag
                                                                    3120
ttaaaagage teagtttgte tegtggttaa atecaagttt tacaaaggte aegetgaetg
laaagtggaa ggtgggctgg ccaggggatc atctagtctg ggtgagaagt gatgataaca
                                                                    3180
```

tgaaggggtg	aagagagatt	tagaagaagt	gattcacagg	attaaacatt	taaataatgg	3240
aagtggagaa	aatggggggg	gcggttccag	atttcaggca	tagatgaaag	aagtgcagtt	3300
aggcacatgt	aaagagaaac	aggaacagca	ggttttaggg	gagaagataa	cagaatgggt	3360
gagaaatgac	acttgagtac	cctagtgtgc	taggtaatca	tctgtctact	tcccttcatt	3420
tgtcatgtat	attcccattt	aatttgcata	aagacttcga	gttaaacggt	cttaccccaa	3480
tttgtcaaat	ttctgcgcat	gatatggtac	aagaaaccgt	aagtggctaa	ggcggcattg	3540
gtgttcaaat	tgcctgacta	caaaggcagt	gcttgttggc	tacattctgt	tgcttcccag	3600
tttagaacat	gttacattga	ggcgcctgct	gcatttccaa	ataaaaaagt	acagaaagaa	3660
ggtggctgta	taaatctggg	gctcacaaag	taattttgat	tactgagagt	ttgctttcaa	3720
ggagcaaact	gtgactcctt	gattatgaac	cttaattt			3758

<211> 3811

<212> DNA

<213> Homo sapiens

```
60
actggaaaac tttgggtgtg agacgggatt caggctgtgg ctaatgtgct ggaagcacgc
                                                                       120
acagligiga ccatcaagta tgcaggaagc aatcattctc ctggctctcc tgggtgccat
gtcaggggga gaagcactac acctaatcct cttacctgct acaggcaatg tggcagagaa
                                                                       180
                                                                       240
ticlecacci gggacticag igcacaagii ticigigaag italcagcai cattgicacc
tgtgatccca ggatttcccc agatagtcaa ctcaaatccc ctcactgaag cttttagggt
                                                                       300
                                                                       360
gaaliggeig teaggeacet acitigaggi igicaecaet gggaiggaac aaciagalii
tgaaacagga ccaaacatat ttgatttgca gatttatgtg aaggatgagg ttggtgtcac
                                                                       420
                                                                       480
agacctgcaa gtcctgactg tccaggtaac agatgtgaac gagccacctc agtttcaagg
caactiggca gaaggictac accictacat agtagaaaga gcaaacccig gaticatiia
                                                                       540
ccaggingag gccincgate cagaagacae aageegaaae atteeetea gitatiieet
                                                                       600
                                                                       660
gattictece ecaaagaget teagaatgte tgetaatgge accetettet ecaeaacaga
                                                                       720
attggactti gaagcaggac acagaagtti ccatctcatc gtggaggtga gggacagtgg
                                                                       780
{\tt aggcctcaaa} \ {\tt gcctccacag} \ {\tt agctccaggt} \ {\tt gaacatcgtg} \ {\tt aacctcaacg} \ {\tt acgaagtccc}
                                                                       840
tegetitace agecegaeae gagtgtaeae agteetggag gaactgagte caggaaceat
                                                                       900
egiggecaat atcacagegg aggateetga tgatgaaggt titteecagee accieeteta
                                                                       960
cagcattacc actgttagca aatatttcat gataaatcag ttgactggta caatccaagt
                                                                      1020
ggcccaaagg atagaccgag atgcaggtga attgagacaa aatcccacca tttccctgga
agtictagtg aaggacagac catatggggg tcaggagaat cgcatccaga taaccttcat
                                                                      1080
```

•						
tgtggaagac	gtcaacgaca	atcctgccac	atgccaaaag	ttcaccttca	gcattatggt	1140
gccggaaaga	acagccaagg	ggacgttgct	tcttgaccta	aacaagttct	gctttgatga	1200
tgacagtgag	gcaccaaaca	acagattcaa	cttcaccatg	ccatctggag	tggggagcgg	1260
cagcagattt	ttacaggatc	cagctggctc	tgggaagatt	gtgctgattg	gtgatctaga	1320
ctacgaaaat	ccaagtaacc	tagcagccgg	caataaatat	acggtgataa	tccaggtgca	1380
ggatgtggcc	ccccttact	ataaaaataa	cgtctacgtt	tatatcctaa	caagcccaga	1440
aaatgagttt	cctctcattt	ttgataggcc	atcctatgta	tttgatgtgt	cagaaagaag	1500
gcccgccaga	acccgagtgg	gacaggtgcg	agccactgat	aaagacctcc	cccagagcag	1560
cctcctgtac	tccatctcca	ctggaggggc	cagcctccag	tatccaaatg	tattttggat	1620
taatcccaag	acaggagaac	tccagctggt	aactaaagtg	gactgtgaaa	caacccccat	1680
ctatattctc	agaatccagg	ccaccaacaa	cgaagacaca	agctctgtca	ctgttactgt	1740
gaacatcctt	gaagaaaatg	atgaaaagcc	aatttgtact	ccaaactctt	atttcctggc	1800
cctcccagtg	gatctgaaag	ttggcacaaa	tattcagaat	ttcaagctga	catgtaccga	1860
ccttgattcc	ageceeagat	ctttccgtta	ttccattggc	ccaggtaacg	tcaacaatca	1920
tttcaccttc	tctcccaatg	ctggttccaa	tgtcacacgc	ctgctgctta	cgtctcgctt	1980
tgactatgct	ggtgggtttg	ataagatctg	ggactacaag	ctacttgtct	acgtaactga	2040
tgacaacttg	atgtctgaca	ggaagaaagc	ggaggctctt	gttgagacag	gaacagtgac	2100
actgagtatt	aaagtcattc	cccacccaac	cactatcatc	accacgaccc	ccaggcccag	2160
ggtcacctat	caggtcctga	ggaaaaaacgt	ttactctcca	tctgcatggt	acgtgccgtt	2220
tgtcatcact	ttgggctcca	tattgcttct	gggtctcctc	gtgtacctgg	tcgtcctatt	2280
ggccaaagcc	atccacagac	actgcccctg	caagactggg	aagaacaagg	aacctctgac	2340
aaagaaagga	gaaacgaaga	ctgcagagag	agacgtcgtg	gtggaaacta	tccagatgaa	2400
cactatcttt	gatggagaag	ccatagatcc	agtgaccggg	gaaacatatg	aattcaactc	2460
aaaaactgga	gccagaaagt	ggaaagatcc	actaacccaa	atgccaaaat	ggaaagagtc	2520
cagccaccag	ggagctgccc	cacgcagagt	cactgctggg	gaagggatgg	ggtcactgag	2580
aagtgccaac	tgggaagaag	atgagctgag	tggcaaagcg	tgggctgagg	atgctggtct	2640
gggttccaga	aatgagggtg	gcaagctggg	caacccaaag	aacagaaatc	cagccttcat	2700
gaacagggct	taccccaaac	cacacccagg	aaagtaaacg	gggtctaagg	aggggcctgt	2760
caatcactga	gatgetgeet	caccctaaat	tctatgggga	tggtgtgggc	atggtgtagg	2820
ggggaaaatg	tgggctgagg	ggattcagac	atccagggtc	aaacatggga	tgtttgacaa	2880
atttttaaac	aaatagaaag	gggtttgatc	acatagttgc	gtgttctgaa	atgatacagg	2940
aacattttct	atcagatttc	agaactacct	gtgcttctga	taagcaagac	tgttaacttt	3000
ggggtgtgga	attgttgtgt	ttcttcfttg	cattgactgc	taggaagctc	tattctgttc	3060
accatagaaa	gtttgtagga	attcctgaca	taaatagtga	agactatcct	tacatctggt	3120
ttccacctta	ttttcctgcc	ctcgttttaa	catcacccag	atttcttcag	ttataaatat	3180
gccatacacc	ttigtaagic	acctcaaatc	ttcttcaaaa	gaagcagaac	agtgaaaaaa	3240

acagatgagt	aagttaagag	ttggtcatct	ggaaagaaga	aaactcagta	ggcaccttct	3300
ttigttttt	cttgtggtgt	ccggatcagc	atcctgcatg	tgagattcat	ccacgttgtc	3360
ctgtctagca	gtagttcagt	tctcttcatg	gttatgtctg	gtttcattct	atgattatat	3420
cacaatttat	ctattctaca	cttgggtggc	agctgcttca	gatttttact	tttaaaaaaat	3480
atacttaaaa	gtgaactaca	ggcagggcat	gatggctcat	gcctgtaatg	ccagcacttt	3540
gccaaggtgg	gcagatcacc	taaggtcagg	agttcaagat	cagcctggcc	tagatggcaa	3600
aaccctgtct	ctactaaaaa	atacaaaaat	tagcttggtg	tggtggtggg	cacatgtaat	3660
cccagctact	tgggaggctg	aggtagggag	aactgcttaa	acctgagagg	tggaggttac	3720
agtgagttga	gattgtgcca	ctgcactcta	gcctgggtga	caaagcaaga	ctccatctca	3780
gaaaaaaaaaa	ataaaagtga	attacaacac	t			3811

<211> 5211

<212> DNA

<213> Homo sapiens

```
ttttagagaa ttttttggaa attaccttta attttatcta agacttctta tatcttaatt
                                                                      60
ttgtgaaaat gtatattgtt cataaaagga aactettatg tteeettact ectaaatace
                                                                     120
taaggagtti teagateeag ttaatgggag attgtaatat teaategtta aaaagtetga
                                                                     180
                                                                     240
tccalacagt attcatttgg tiltttaaaa agtitticaa agtattigit tigaggaaag
aatgcaaltg gatatttaat giggtaaaat ittigcaaaga tiattictit itagttagaa
                                                                     300
                                                                     360
gagtglaatt aanagtatta atticttacc ticcacacgc gigcacagcg ganattiigt
gtiliticcii titictiilag cagiccatti igittaacac acagatccca aalittigaga
                                                                     420
ataaatatgi cataaagaaa tagggtatci tcaataccii tggtataagg gitaatcaca
                                                                     480
gtttatttcc caaagtgaca aactggacac aggttaaata agctgttaga gtggtaacat
                                                                     540
                                                                     600
tgtaatgcat cagtacttta gaatatggtg caggcattaa aatccctggt ttcagagaat
cttcagtgac ctggtaaatg tttacatgtc aattaaagaa gcacatgaga ctgaatgttg
                                                                     660
                                                                     720
tataatetea titileagaaa aaagiitgig catatagaaa igigictaat aaaegeaaaa
                                                                     780
ggaaagtaca totgagtact aacaacggat ttgagcggga ttattgatag attattttc
                                                                     840
telliatati etgiallila aaaggigtaa eagggaleea caltititat glagtitaga
gggaaatigt litaatiitt giteateige itaeetilei aattiigiag teaggeetii
                                                                     900
                                                                     960
ctactttgct gcctctttaa accaaacgta ataaacttgg agctgtcact gtatgccagc
atcataaaca ccatcatttt atgataggga aaattttttg gctcacttgt ttagaaaatt
                                                                    1020
agtaaaatti attagcatta tiattiatta galtigitto ticattiigi tagtaigota
                                                                    1080
```

```
caatttagca tctttgaaca ttatacagaa tgttgacttt gcttaagggt tgtttgaata
                                                                    1140
ggcatttcaa agtgcttitg ctttiggctg catggagagt agaatctatt gaggtgattg
                                                                    1200
ttettgtgat giggtgecat giteeaaaat taatatatat geatggtatt aatgaggaat
                                                                    1260
atgitigeat teataittia geagataeaa titateagig tiggigaeaa eetetaiggi
                                                                    1320
tttattttct ttataataca gtcttttgcc tggatggagt cctcacttta aggttaagag
                                                                    1380
taactaagcc aatgttactc cagctacagt tccctaaatt atactatagc tgctgggaac
                                                                    1440
aaagccatgc tgatgaatct ggacttgtgc atgatttttg tttgcttctc attaacctgc
                                                                    1500
ccacceteca etecaaaatt ataceteatt aacgttetga taacageeag gaagacagee
                                                                    1560
                                                                    1620
tcacctgaac cctctttgac tgaatggatt tttcattgtt tttcttaaat gcctacgctt
cagaggctat caactgctta aatgcagcca tcgacattta cacagacatg gtaagacatt
                                                                    1680
geattgettg agtggetgtg gggtggagte ttgagatgge ttagagttet atetttettt
                                                                    1740
                                                                    1800
ttlatgitcc caaaciggca ticagatagg laaaatcggi gigigacigi ticiigiiit
ttcccctagg gaaggtttac aattgcagcc aagcaccaca ttactattgc agagatetat
                                                                    1860
gagactgaac tigtagacat igagaagget attgeacatt atgaacaate igetgattat
                                                                    1920
tacaaaggag aagaatccaa cagctcagca aacaagtgtc tgctgaaggt ggcagcatat
                                                                    1980
gctgcccagc ttgagcagta ccagaaagcc attgagatct atgagcaggt tggggcaaac
                                                                    2040
acaatggata atcettigit gaaalacagi gcaaaggati acticitcaa agcigcecte
                                                                    2100
                                                                    2160
tgccacttca tagtagacga gttgaatgtc aagcttgctc ttgagaaata tgaggaaatg
                                                                    2220
tttccagcat ttactgattc aagagaatgt aaattattga aaaaactcct agaagctcat
gaagaacaga acagtgaagc ttacactgaa gcagtgaagg aatttgactc aatatctcgc
                                                                    2280
tiggatcagi ggcigaccac caigiigcii cgcaicaaaa agiccaicca aggggaigga
                                                                    2340
gaaggagatg gagacctaaa algaaatgit titgicittg iggcaigcag claactecic
                                                                    2400
tttagttiig iettagggie aagigaieti talgggaige etaittaaig getiaaitti
                                                                    2460
gitgcatatg agecagaegg cetgigtati gittaagete gecaagleig igitgeigtg
                                                                    2520
aaatgaatga aggagagget eetgtteate tigtggtaat gatgggtigt ticatgetta
                                                                    2580
teagaacccc cagegiilte igagaagiae ileagaalei calleeleat allicatigg
                                                                    2640
tattigtgga gcctatgiti aaigitgcca cgigiitita igicciiiti giiggaciig
                                                                    2700
agtactcage ccagttgite teatggatge ittgcattit etetgtgeit tggcatetga
                                                                    2760
                                                                    2820
atatgiicit taaaigigig tilagiitag gacagilaci aggaalgagi tialaactic
                                                                    2880
attagaaate atticiatit ligilaicel gigallaiti igalggigel agigaciagi
                                                                    2940
ticitigeti ittigigilgi teegiatget aacatgigea iggeaaaaat itagaatage
cagggtetgt aggeateaca tigigaggaa gggagetite iggaagtaet igeticatgt
                                                                    3000
atgatgagig tcaaagigaa tilgatiigi acitagacac acgcgiitac acacacac
                                                                    3060
atatcacaag atcigitaga aaiggaatii iicicittii ciggagatag iiticactii
                                                                    3120
tagttggagt ggaaatccct tlatatttac attgaagtat tttaattggc atagcctgct
                                                                    3180
cattattite atgittatae actileceae gitgaggigg igigiletgi getgigaeta
                                                                    3240
```

```
3300
tagaaatett ggteaggget ggatagatta tetaagteaa gettgagaat gaatgtatgt
aattileetg titatigtae atgatgggti aggtggggtg aatgiggtae aggaalgtee
                                                                   3360
                                                                   3420
tgtatgccca agtgggcaag aaccccaact tgtttetcag gggacttgat tgttctctta
gctggtggaa tatgttggct tatgtgtttg aactctgtcg tgittaattg gtttatataa
                                                                   3480
                                                                   3540
tatatgtatg ctatcttgat tcatgaactt gatcctatta atttatatgc tgatattgta
                                                                    3600
ctttagacat acgctigtet cctgaatgic cictgaatat tlatagitaa atgattiata
                                                                   3660
ttlgaaatgt gttgccagac ttaacccagc agacactctg acatcacgga gcttcactga
tgacaggtaa cgaaacttcc tatgttatgt caggtagtag taagtagtat tggaatgatg
                                                                   3720
                                                                   3780
ttttcatttt tggtggctct caactggaat tggtagtgtt tccaggccaa gggtcgactg
                                                                    3840
caggitgitt gagaaatgat gagtaggica giclaggaag aaagagaaag tagcaggaaa
                                                                   3900
ggaagtggga agggccagcc aaggacagac tgtagaggat ccacatcagg tggccacgag
                                                                   3960
gactigcagg ctatagttat ggtggtgaca igcatgaggi gggciggiag agcaggaagc
                                                                   4020
tetgigatgi cagagcatet aetgggacia caggigeaci giagicecca etaciggggg
                                                                   4080
tggcaatgaa gacactetgt etgttgggee etagaattta atgtggattt eeteetteet
                                                                   4140
tecaagitet gagattetta aatgagaget ggetglette tagaggtaag acetggaatg
                                                                   4200
gagtecagtt ggtacttttt cactecetet tagaatetet tatgaaaaaa tgateagaga
                                                                   4260
gaaaagtggg gttttgtttc cccacctaat aatatatect acaaccagec aaatgcactt
                                                                   4320
ttgtgaaaat ggggtgtgag gagtggttct gcagctlgag tcctctggtt ttaagtagtt
                                                                   4380
tgtttctact tgtttaaaga atcttctggt ctgaccactt aaagtaaaaa ctacatgatt
                                                                    4440
tattttcggg caattatgtt tagetttcat cattatactc caacagaccc gtclgaaggg
                                                                    4500
gtattitttt ttaaaataat gtttgtaaca tittgtigtg tcaattagag ggtcacttgt
                                                                    4560
tigiatigca ataaacactg ggaccagtic cggggilaag aatlaalitt tgiltilaat
                                                                    4620
atticacatg aaaagaatca aagtaattgi aatggclaga agagaccigc cagaagatta
                                                                   4680
aaaaaaagaa tgagagaaaa gcccagttag tggtgtgcaa acttacttcc tttaaatgtc
                                                                   4740
ccatggatgt aggacagtgc catgtttcaa gatgcctgtg aactaggtct tcaagattta
                                                                    4800
tagaatgtta cttatgaaca aaatataatt atttatggta caattettgt actttagcaa
                                                                    4860
atolggagti agticatagt caaagtoagt taatatiici tagaggaaag titigotiti
                                                                    4920
tgtggcaaca tttttatagc ttgtgtgagt tcttttttat ttaatgattt gaaagcagta
tttttgcaca gtcgtgaccg tgtgtggtgg catcactgta accaaagtat atgcaccagc
                                                                    4980
                                                                    5040
cettgtgcat ttattgttte teetgatttt gtggatttaa atgteeaaat geaaacettt
                                                                    5100
gigacticci tiggaggaet iggeageaea geaigeeece gigaeeigee igeigiggta
                                                                    5160
tgagctatga ccaagagcag gcttcctgct ccatggagtc ctgagttgct ctggggcagg
                                                                    5211
ggattacgtt atgaaaacta accatgtgta acaataaate taccttagca g
```

<211> 3722 <212> DNA

<213> Homo sapiens

<400> 2038

60 agacttgatg ttttatatag aaatggaccc accaggtaat actgcagtat tattgtagag 120 agttagttaa titcgtggct tittaatiit tcgaaagcta cigtaaaaga tcctiitigg atttctgttt ttattaattt gtttcattga taaaaattag tttgctcatg gcttaaaaat 180 240 taaacagatt gtttgactgt ctgtggaagc aagcagctca ggctgtgtgt ggtaaatgct 300 tattettaet tgaatggata tgaattgaac teeagttttt caetggtgte ttttgttaat cgagatcctt ccctgggtga gttatgttgt gggatattgt ccctgtaatt aaaatgatgc 360 420 atcttttgtg ctgcttttct ctgttgccag tggatgagaa cagtgtagca ctttgcagtg ataacacttg gtactttaga aagcatgtaa aatgtagcag tgattacaac teagttetet 480 540 aaatgttgag acttlgcttg ctctctcata ttaagatatt ataatgaaaa aagaagttga ctttccatta ttgttagtct ttgtaaaata ttcttggtag atacctgaaa tcattttttg 600 tataagttaa aatagtaaca gtgctttaaa acttatgaca gaatttacct aaaaatccta 660 720 gatttatttt gtttcctaag taagttgttt tattccaatg ttagctctcc ccctgccccc 780 atttaaggta ttcaggaata ctgcagtctt ttatttgtca ccaattggta tatatgaata 840 ctgatttgac attgaggaag ggggatgtca tttttaatca gacctagtat atagagcaca 900 atttatccaa cagaatatta acatattaaa gagatttagg gcacagatga gagtttctta 960 aagtggcttt tggcagaaca gtgcctgaaa tactaagatt agagaaaccc aattgctcct 1020 cttaaaacat actgctgtag atgagccttt ttattactgc aacagagttt gtggaggaca gagaccaaat tigiciticg taattaaata agaggaaatt aaagccaact catgitatic 1080 ctgctactca tatgttcata gtttcttact ttagatggat ttgaccagge atgaaacttt 1140 aatataacta gaatctagaa gtacagaatg tcatgactct ggatttactt tgaaatttat 1200 1260 teacatggcc ageccaattt atttgttagt ttetaagget etetetitt teteettite agtttcattt ctttttgagc catgctctga aagatttttt ttaagaaaat tatcttccat 1320 attgcatgga attgtgaact aatgctatat atttcagtta cictaacttt tiattititi 1380 1440 aaagtaaaag tattcatcta aagaaattta gttctaatgt agttgggatt gcgaacaact 1500 ttitetttt catelgeage aelgeeleel aaaceaceaa aacelaelae iglageeaae 1560 aacggtatga ataacaatat gtccttacaa gatgctgaat ggtactgggg agatatctcg 1620 agggaagaag tgaatgaaaa acttcgagat acagcagacg ggacctttti ggtacgagat gegtetacta aaatgeatgg tgattatact ettacactaa ggaaaggggg aaataacaaa 1680 1740 ttaatcaaaa tatticatcg agatgggaaa tatggettet etgaccealt aacetteagt 1800 totgtggttg aattaataaa coactacogg aatgaatoto tagotoagta taatoocaaa 1860 ttggatgtga aattacttta teeagtatee aaataceaac aggateaagt tgteaaagaa

gataatattg	aagctgtagg	gaaaaaatta	catgaatata	acactcagtt	tcaagaaaaaa	1920
agtcgagaat	atgatagatt	atatgaagaa	tatacccgca	catcccagga	aatccaaatg	1980
aaaaggacag	ctattgaagc	atttaatgaa	accataaaaa	tatttgaaga	acagtgccag	2040
acccaagagc	ggtacagcaa	agaatacata	gaaaagttta	aacgtgaagg	caatgagaaa	2100
gaaatacaaa	ggattatgca	taattatgat	aagttgaagt	ctcgaatcag	tgaaattatt	2160
gacagtagaa	gaagattgga	agaagacttg	aagaagcagg	cagctgagta	tcgagaaatt	2220
gacaaacgta	tgaacagcat	taaaccagac	cttatccagc	tgagaaagac	gagagaccaa	2280
tacttgatgt	ggttgactca	aaaaggtgtt	cggcaaaaga	agttgaacga	gtggttgggc	2340
aatgaaaaca	ctgaagacca	atattcactg	gtggaagatg	atgaagattt	gccccatcat	2400
gatgagaaga	catggaatgt	tggaagcagc	aaccgaaaca	aagctgaaaa	cctgttgcga	2460
gggaagcgag	atggcacttt	tcttgtccgg	gagagcagta	aacagggctg	ctatgcctgc	2520
tctgtagtgg	tggacggcga	agtaaagcat	tgtgtcataa	acaaaacagc	aactggctat	2580
ggctttgccg	agccctataa	cttgtacagc	tctctgaaag	aactggtgct	acattaccaa	2640
cacacctccc	ttgtgcagca	caacgactcc	ctcaatgtca	cactagccta	cccagtatat	2700
gcacagcaga	ggcgatgaag	cgcttactct	ttgatccttc	tcctgaagtt	cagccaccct	2760
gaggcctctg	gaaagcaaag	ggctcctctc	cagtctgatc	tgtgaattga	gctgcagaaa	2820
cgaagccatc	tttctttgga	tgggactaga	gctttctttc	acaaaaaaga	agtaggggaa	2880
gacatgcagc	ctaaggctgt	atgatgacca	cacgttccta	agctggagtg	cttatccctt	2940
cttttcttt	ttttctttgg	tttaatttaa	agccacaacc	acatacaaca	caaagagaaa	3000
aagaaatgca	aaaatctctg	cgtgcaggga	caaagaggcc	tttaaccatg	gtgcttgtta	3060
atgctttctg	aagctttacc	agctgaaagt	tgggactctg	gagagcggag	gagagagagg	3120
cagaagaacc	ctggcctgag	aaggtttggt	ccagcctggt	ttagcctgga	tgttgctgtg	3180
cacggtggac	ccagacacat	cgcactgtgg	attatttcat	tttgtaacaa	atgaacgata	3240
tgtagcagaa	aggcacgtcc	actcacaagg	gacgctttgg	gagaatgtca	gttcatgtat	3300
gttcagaaga	aattctgtca	tagaaagtgc	cagaaagtgt	ttaacttgtc	aaaaaacaaa	3360
aacccagcaa	cagaaaaatg	gagtttggaa	aacaggactt	aaaatgacat	tcagtatata	3420
aaatatgtac	ataatattgg	atgactaact	atcaaataga	tggatttgta	tcaataccaa	3480
atagcttctg	ttttgttttg	ctgaaggcta	aattcacagc	gctatgcaat	tcttaatttt	3540
cattaagttg	ttatttcagt	tttaaatgta	ccttcagaat	aagcttcccc	accccagttt	3600
tigitgcttg	aaaatattgt	tgtcccggat	ttttgttaat	attcattttt	gttatccttt	3660
tttaaaagta	aatgtacagg	atgccagtaa	aaaaaaaaaa	tggcttcaga	attaaaacta	3720
tg						3722

<211> 4323

<212> DNA

<213> Homo sapiens

60	gaaataatta	gcgaaaagga	ccctgctgtg	cttgacactt	gctcaggttt	acagggagtg
120	gcccagctga	cactgcaagc	gcgtcggggg	atcgctgatc	gggctctagg	acagctcctg
180	ggacctcccc	cgcgggcctc	cctgcgcccg	aggcgccgcc	gggaggagac	gccatgctct
240	ggctcaccag	tgactatgat	gaggaaaaga	tctcaaaaaag	cctcagacca	tacaggatat
300	acttaattct	ccagggaaca	acattcataa	taatgagaag	ggatcttcat	tacccctctt
360	tttatatgga	atctgcatat	aaccagaaaa	tttttatgag	cctataacat	ttattgaaga
420	gggagtaaaa	acattttctg	ggaatgctgg	aattgttgaa	atggcaaact	gagactgaag
480	tagtatgaag	cttcttttac	aagccattct	acaagatgag	agctaaaaat	cgacctatac
540	cgatctctat	gggaatttga	acacgctggg	caaaggaatg	tcttctccag	tcatcagacg
600	ttcccaggaa	aaaaaaggaa	cctatgtctg	gacccagatt	agctggacag	cgtattagtg
660	accagactcc	caaaggatga	aagccacatg	caacaccctg	cttatcacag	gactatttat
720	gaagcctctg	gaaaaaggat	gctctggtga	gagtgaagca	atagaaccat	ccagtgctct
780	cttctataac	ttaatggaca	agagcctcta	acagaaaaaat	gaaaagaaag	atgatggaca
840	agtaaacagt	ctaaggtcag	gaatcagaaa	tccagccttt	caattttcat	catgaaacat
900	tgaaaatagt	aatttaagat	cttctccaaa	aataaagcaa	ctgaagaagt	aacatgagaa
960	actaaagaag	aacaaagacg	gcaacaggag	cattattttt	ttgctcttca	ccccaggatt
1020	tgctcgcatt	ctgaaaagaa	cagggacctt	gaggctccta	cgctactgca	acagacattc
1080	cattaacttt	tggctcagta	agcagtgatg	agaagaaatt	ataaagatgc	ttcctcatgg
1140	aagagagatt	aagaagagaa	agattaaatg	cattcttcaa	tcttggaatc	cacttttctc
1200	tcttcaaaat	tactgaaatg	aaggcgatta	caataaagaa	taacaaaatt	caaagaatag
1260	gctaaaacat	agcttctatt	tagcagtaca	gacaacagtt	taaaaacaga	aaactagtaa
1320	ttgcatttat	agttctgtac	tgaatgcata	tactctttga	tcagagatat	ttcaaaaaaac
1380	tcatttcttt	caaaaaaagc	aattgaatgt	aatcgtagaa	atgagacttg	acgaacatat
1440	gtttagtttc	atgagtttga	gttggacaaa	agggttcaca	gaggttaatt	ttgaagtgat
1500	aaaccatgta	cttttatagt	tgccaaatag	atactgcata	ataagcttga	agtaactgaa
1560	tagcttggac	aaacttcatt	taagcagttt	tgtcttcaga	atttaaatgg	atgaactcaa
1620	aagaagctgc	actcagaaaa	ticagaaatg	aatcaatgga	actgaaacat	tctcaagaga
1680	cctgatagtg	aaggaateta	ttacaccatc	aaatacagtc	aatgaaaaag	cagttcttgg
1740	taaatggcct	aggactatct	cataaaatct	ctggcatttt	cttgaaaaact	acagtagctt
1800	aaaaattatg	ttiggaaagg	ctgtctggcc	aacatcagag	ggctatctgt	gttgacttct
1860	ttttattctc	ccccagccc	tctgaacctc	attgaaattt	agaaatccta	gactctgtta

tctcttctgc	tgatgaaaga	cctttcatca	gttcaaagct	tttcttaagc	tctttttaa	1920
gttaattgaa	cttttcttt	atttattttt	caaaaaaaatg	tttatatcac	atagacatat	1980
tacatcggct	aaagcaagac	ttggcccaca	aatacctatt	tgttgctgaa	tgaatacaat	2040
ggataaagca	aggctgttgt	agctgaagtt	acatagggaa	tcccaaactc	tgccctctta	2100
gcatcttatt	ctacatgaca	actctcaagg	tactcacaga	tctgtttaac	ccacttgaaa	2160
aaaaaacact	aaaaatgaag	aaatgctata	agtataaact	atgattttat	ttataaattc	2220
tgtattaaaa	tggaattata	tgcaacattc	tttcattctg	taaactaatt	ccatttgcat	2280
tcctcataag	cattgtagta	aattgatcat	attacatgta	ctaaggaatg	agattatatg	2340
cagtaaaccc	aactggaaga	ttaacaatat	taaaatatga	aacattttta	agacaaaggc	2400
attacttctc	agtattacca	aacctaaact	ggttgaaggt	gaaagtgtgc	tatggccttt	2460
tcaagcctaa	gaagtctctc	ttactgagta	aaccagaggc	ttgcatcgct	attctttcac	2520
ctgtcaatat	taataagaaa	atagtctcat	ctcacttaaa	tgaggcaaat	gtaatagtta	2580
aaattcaaca	tacttataaa	aaactagtgt	catgtacctg	ccatgaacat	gacaaaaggt	2640
tagtcttcaa	tagactgaaa	tgtataagag	aagaaccaag	tcttacatag	aaaaaaaagg	2700
tagatatgaa	aagaaaaaatc	acagaagaga	gaatgcaaat	ggccactaag	tatatgaaaa	2760
aagtcgtatc	ttaacagtga	acaactgtgt	tagtctgtat	caatcagaag	acagaaacaa	2820
ggtagtaatt	taaacaggga	aagtttaata	taaataataa	ttaagctatg	ataggagaat	2880
aataataaag	atgaaaagag	aaggtaccct	aaggctgagg	gaaagaatcc	taacaaggaa	2940
aggcaggaat.	gagggtttca	gaattcactg	gagaaggtgt	ggttgcagcc	cactggagag	3000
aagtttgctg	gcttgcccag	gccagagcag	gaccacagat	actggacaag	ctggtacagc	3060
caacccccta	ggtgtggacc	agctgaggca	ggtgggcaga	tatgcagagg	gacttggggc	3120
tttgccaaag	ggtaagcaca	aagaaggagt	cacgggttct	gttcgaggca	ctgttgggat	3180
taggagtcgg	agggacctac	tttgcaggaa	cctagcataa	ctttgtgtga	cgagactgca	3240
caagaca a ag	ctcaggcaag	tggctcagta	gttggccagc	ccagcagggt	cctctgtatg	3300
agtgtgcacc	cagctgaaga	gaagaaatgg	agagcagcaa	ttggagcttc	aggaccggct	3360
tgcactgtgg	ctccaggita	taccaccact	gcccaaagca	aaagctagag	aagcaagtgg	3420
agaaatgctg	gagaaagctg	caccctacag	gcaaccagca	ctgcagaaac	cactccaggc	3480
aaagtagtga	aggaaaaaaag	cctgctctcc	agtagcctgg	cctgtcagcc	tggaggaatc	3540
aggaaagacc	ccttcctctt	gcagtgtgtc	tccagcgccc	tctactgaca	aagtatgcca	3600
tcatgcaagc	tgcaaaggaa	acatttcaag	agtctatatc	tattttcacg	gagcgggcaa	3660
ccaacagtga	atgtggagct	gagagacagt	aaaataataa	ctgacatgcc	accgaagtac	3720
aaagtaaaat	aaataaataa	atacacattt	tggcctatta	gcaaagatta	agaaatgata	3780
acattaaata	ctcaataaat	caccatgaga	tggggactca	aacttctggt	aaaaatacaa	3840
atagatataa	tttttcttga	aggcaattta	gtagtctgtt	tatcctataa	ttctacttgt	3900
aagatcctat	catatgaaaa	taaccagaga	tacaaagaca	ttctgcaaag	atatgtttta	3960
tattgttatt	tattgtgaca	aaaggaaata	aaaagcctaa	atgttcagaa	aattattta	4020

aaagatgaaa	gagggaaata	ggccatggac	ggtggctcac	gcctgtaatc	ccagcacttt	4080
aggaggccaa	ggtaggtgga	tcacttgagg	ttgggatttc	gagaccagcc	tggccaacat	4140
ggtgaaaccc	tgcctctact	aaaaatacaa	aaaatgagcc	gggtgcaatg	gcaggcgcct	4200
ataatgccag	ctactcggga	ggctgaggca	ggagaatcgc	ttgaacccgg	gaggcggagg	4260
ttgcatgagc	cgagatggcg	ccactgctct	ccagcctgga	cgacagagca	agactccgtc	4320
tct						4323

<211> 3646

<212> DNA

<213> Homo sapiens

<400> 2040

taggetgtet gactaggggt acaggatetg tgtagtaaac acttggaaga etcagtgtte 60 ttateaaggt cagetaatee tgaactttga eeetteeett aggeattget ggatgteagt 120 180 aactaagcat gaatttaggg tcgtagctgc ttttgaccca ggttggagga ttgccagggg 240 ccacctggga agggctgtgg ttctcacctg tgctctgagc tcctcttgca gagttccagg etggaccetg eccagecate eccettacce tetgeettet tggtacacag acceccaaat 300 360 gacaatgcaa gtcagagaat ggtgtaaaag ccgtggagtg gagtcaggag ctgagttcct 420 gteceeatgg gttetteaag aaaacaggte attggeetgg atgataeetg aggggtetet 480 ggccctgact ttttctagtt gaaagaagag aatgccctca actgtccagg gctctgtgtt ttecaccaga etcatteate cateaaagae ectecageee atetteacag accectettt 540 tetectiett teeteeteae tieteeteet eeetitigit taleigieet ateetiteet 600 cacttectga geagagattt etgtaaaaat aaatgeacat ggeeetgget tgtaeagete 660 720 acagattage aggetgggae ggeeaggaee ceagggaeee tggtgggaag tatacaagge 780 tggatggcc ctggatggac gagggcaggg aaagccggcc agaagtttcc tgaggtgctg 840 acagtgatga gaagcccaca gggcagctgc attgctttgg ccttctccgg acccacagcc 900 ctctctcagg ctcccatcag cccaagttag cagctacctc tgagctcacc cacgggaatc ccacccctc ccagagtgac aaattttaag ctaagaagag ggaaaggact tgggtggaga 960 1020 aaaccaagtg tecagtetga ettgteacag ecaaagcaca gegetgeagg acatggetat teeceegae acageetetg acceeteeae aaggeatgaa tigaggiegg gggaggeagg 1080. 1140 caagcaggec agaccatagg cagctgatgc agggactgga gaggcaagaa gccgatgctg 1200 agctagaagc cttctgtgga acaggctgga ccccagatgg cctgggatgc gggggcctgg gttgagcggc gggggccaca ggctgctgct gtactgccca ttggacacac ggttcagggt 1260 1320 gcctcaaaag ccactaaaca cacgcctcaa ccttctggtt gtctgtggct taccacttgc

ctggaaacat	tcactctagg	tcacatgatc	ttcctcccaa	cccaccctct	tcctcctcct	1380
tctgggaggt	gccaacagag	agccccctgg	gagcctgggc	tgctggtgga	agcctggctg	1440
gaggggagag	tctccctaga	gtggactgac	gcgctgccac	ctctgcaaag	cctcacagcg	1500
gccgcccctt	cacagatgca	gaactgaggc	ccagagagcc	ggggactagg	aggtatcaag	1560
tccaaggtcc	agccaagatg	tcctgcctgc	aggctgcctc	ccagctgcag	gcctgcaagg	1620
tggggtgctg	ggggtgtgga	gggcgaaggt	ggcacgggtg	caccagcagc	ccttctgggc	1680
caaaatacac	ctgacctgcc	tgtacagcac	cccaagtccc	cttgcttaac	ctgggtcccc	1740
cttttctctg	aaaaatatga	gacttgtttg	gtccttcctt	cgtttatcct	ttctttttt	1800
catttatcaa	atgcatgtta	agctctcgct	agtgccacac	cctgtgcaag	agatggtgag	1860
gatgataaaa	tgatgatatg	ctatcatgtc	atcaaggagc	ttaagtctaa	taatactaat	1920
actaataata	acttactgaa	tgtttattac	atgcccggga	ttgtgctgca	tgtactacct	1980
catttaaatt	tcaaaacaat	cctatgagat	ggaggaacta	ttcttatccg	catttggcag	2040
agaaggaaac	tggagctctg	agaggggatg	tgacttgcca	gggctgcaaa	gcaggcaggc	2100
aggatgaggg	ttctcatcag	gcgtctggct	cagageetet	tggggagaca	gacgcacagc	2160
acagccctga	ggcctcttgc	cctagcacgt	tatgcttaat	gtatgtcaaa	atcaccctct	2220
ttatcttaca	gatgagcaaa	ctgaggccta	cgcaaagtca	cggctagttt	gcagttgtgt	2280
cagaccccag	cgctgtggtt	ctgatgccag	cttttacctc	tggccttcag	tttcctcttg	2340
cttgcctgaa	cctaggcagt	ttccttagat	gatccccaag	ttctgaaatt	ctgattgtat	2400
gatgttagcc	taagacatgt	tagggagaca	gaacagagag	gcaggaatgg	ctcagctgaa	2460
actagacctg	gagccctgcc	acatccacaa	gcaccccggg	gaacaatcct	tgcccagtag	2520
ggagttaaga	atgttgaaat	geggeeagat	gcatggctta	tgcctgtaat	cccaacactt	2580
gggagaccaa	ggctggtgga	ttgcttgagg	ccaggaactc	aagaccagcc	tggccaacat	2640
gatgaaaccc	tgtctctact	aaaaacacaa	aaattaccca	ggcgtggtgg	catgcacctg	2700
taatcccagc	tacttgggag	gctgaggcag	gagaattgct	tgaacccagg	aggcagaggt	2760
tgtagtgagc	caagatagtg	ccactgcact	ccagcctggg	cgacacagag	agactcagtc	2820
tcaaaaaaaa	gaaagtggaa	atgttttctt	gcttcaaggc	acgtgacttt	taactcaatt	2880
gaagaaaagt	atgcgtgtat	tgatagagat	ggccatcaga	ggaactgaca	ggtcttagca	2940
gttacagatg	agtttcctct	agaggtcagg	gaagagggag	aagatacaaa	gttctttaac	3000
ttacagtctg	aggcaaaggt	gaact taaca	gggccagcaa	gatccttaca	tggtgaggta	3060
agagggccca	aatcagccaa	getgeeacit	ctgcagagcc	cgtgcccttc	tccacctgtg	3120
tcggtggagg	ctatcagcct	cagccccttg	tctgagttat	catagcctcg	ctagcatctg	3180
tctcagcccc	aacccttcca	aaagccaggg	tgacccattc	agctactcct	ttgcgaggaa	3240
gtgacagcag	cctggctggg	ttgtgggtgg	gggagtggtt	gggggtctct	gttgccctgg	3300
aaggaattcc	tacagtaagc	ctgagagete	ciggccaagt	gtggctacag	aaaggaacaa	3360
aatttggggg	gctgagggca	agagagggag	aggattaggg	atgctgctca	gtttctcttg	3420
ataaatggat	cctgctgcct	gaaggatggg	gagctcccag	agttgggtgg	agccatgaat	3480

gggccaccca ggacgtgga gtgagtagta agaaaagggg gaaggaggtc aggtgcggtg 3540 gctcacgcct gtaatcccaa cactttggga ggccgaggtg ggcgggtcac ttgaggtcag 3600 gagttcgaaa ccagtgtggc caatatgctg aaaccctgtc tctatt 3646

<210> 2041

<211> 3679

<212> DNA

<213> Homo sapiens

<400> 2041

attgctgtgt caagttccag agaaaagctt ctgttcgtcc aagttactaa ccaggctaaa 60 ccacatagac gtgaaggaag gggctagaag gaagggagtg ccccactgtt gatggggtaa 120 180 gaggateetg tactgagaag ligaccagag agggieteac catgegeaca giteetietg tacctgtgtg gaggaaaagt actgagtgaa gggcagaaaa agagaaaaca gaaatgctct 240 300 gcccttggag aactgctaac ctagggctac tgttgatilt gactatette ttagtggccg cttcaagcag tttatgtatg gatgaaaaac agattacaca gaactactcg aaagtactcg 360 cagaagttaa cacttcatgg cctgtaaaga tggctacaaa tgctgtgctt tgttgccctc 420 ctatcgcatt aagaaatttg atcataataa catgggaaat aatcctgaga ggccagcctt 480 cctgcacaaa agcctacagg aaagaaacaa atgagaccaa ggaaaccaac tgtactgatg 540 agagaataac ctgggtctcc agacctgatc agaattcgga ccttcagatt cgtccagtgg 600 ccatcactca tgacgggtat tacagatgca taatggtaac acctgatggg aatttccatc 660 giggatatea cetecaagig ilagitaeae eigaagigae eeigiiteaa aacaggaata 720 780 gaactgcagt atgcaaggca gttgcaggga agccagctgc gcagatctcc tggatcccag agggcgattg tgccactaag caagaatact ggagcaatgg cacagtgact gttaagagta 840 catgccactg ggaggiccac aaigtgicta ccgtgacctg ccacgictcc catitgactg 900 gcaacaagag totgtacata gagotactic otgitocagg igccaaaaaa toagcaaaat 960 1020 tatatattcc atatatcatc citaciatta tiatiitigac catcgiggga ticatiiggi tgtigaaagi caaiggcigc agaaaatata aatigaalaa aacagaatci aciccagtig 1080 tigaggagga igaaaigcag ceciaigcea gelacacaga gaagaacaal ecicletaig -1140 atactacaaa caaggtgaag gcatctcagg cattacaaag tgaagttgac acagacctcc 1200 1260 atactttata agtigtigga cictagiace aagaaacaac aacaaacgag atacattata attactgtct gattitctta cagtictaga atgaagacti atatigaaat taggttitcc 1320 1380 aaggitetta gaagacatti taatggatte teatteatae eetigiataa tiggaattit tgattettag etgetaceag etagttetet gaagaaetga tgttattaca aagaaaatae 1440 atgcccatga ccaaatattc aaattgtgca ggacagtaaa taatgaaaac caaatttcct 1500

```
caagaaataa ctgaagaagg agcaagtgtg aacagtttct tgtgtatcct ttcagaatat
                                                                    1560
tttaatgtac atatgacatg tgtatatgcc tatggtatat gtgtcaattt atgtgtcccc
                                                                    1620
ttacatatac atgcacatat ctttgtcaag gcaccagtgg gaacaataca ctgcattact
                                                                    1680
                                                                    1740
gttctataca tatgaaaacc taataatata agtcttagag atcattttat atcatgacaa
gtagagctac ctcattcttt ttaatggtta tataaaattc cattgtatag ttatatcatt
                                                                    1800
                                                                    1860
atttaattaa aaacaaccct aatgatggat atttagattc ttttaagttt tgtttatttc
ttttaagttt tgtttgtggt ataaacaata ccacatagaa tgtttcttgt gcatatatct
                                                                    1920
ctttgttttt gagtatatct gtaggataac tttcttgagt ggaattgtca ggtcaaaggg
                                                                    1980
tttgtgcatt ttactattga tatatatgtt aaattgtgtc aaatatatat gtcaaattcc
                                                                    2040
                                                                    2100
ctccaacatt gtttaaatgt gcctttccct aaatttctat tttaataact gtactattcc
tgcttctaca gttgccactt tctcttttta atcaaccaga ttaaatatga tgtgagatta
                                                                    2160
taataagaat tatactaitt aataaaaatg gatttatatt titggicatg titgtaagag
                                                                    2220
agtgaatgca cgtgtgagaa cattagcttc ttctgaactc attatatctc cacagaggtg
                                                                    2280
ttgatacitg atgcctaaca gittigcaga igigctacat iggaatigig tattittatg
                                                                    2340
glglacattc tattgtgata tatttattga ataattaatg tctattgacc atataagtgg
                                                                    2400
cgaaaaatgc accatagagg acatggggta tttatttaca aactatgagc tacataataa
                                                                    2460
gcaagtggcc atgggatggc atgaccetec cetecatatt tttgtggagc aaaatattgg
                                                                    2520
                                                                    2580
caatgtttat gtaaatcatt gttaatatca tgaaattatt tttaattaaa aacataagtc
                                                                    2640
tatttgctcc atagcagaaa aaacatgaga agttttttca tcatgataga aattgaaaca
aactatattc attetteaat cataceatet gagattttta agacagetet tttgtettat
                                                                    2700
                                                                    2760
aagtatattt ttctccctct agacatttca gttactatgg attttgtcct caaagggact
tttagtetat tttggatgta aagetaatet aatgacaett ggeacatgat attttgatea
                                                                    2820
                                                                    2880
agccattitg actigaccaa aaagcagtgt ccattaggit tctgcatata aatattacca
agcaatgitc acaatagaca tcattacact giccitgaaa titattaatt ciicatecaa
                                                                    2940
ccctggttga gctgaggctc atagttaggt tcaagactat ctgtttaaat attactgaaa
                                                                    3000
                                                                    3060
aacaaagtaa gacagtacta tgcttacctc ttaacttgat aatgtcaaaa caggcatgtt
aaatgacatc atagaaaaga ciicaagata attitatagaa gitaaattat atigtacaga
                                                                    3120
                                                                    3180
aaataattgt atgaaaatct ctactatggg gctggaacat ggttgaacat tagaatgata
                                                                    3240
taaaaaatta tatatattet eeaaateeac getagaeetg teaaattaga gaatetagag
                                                                    3300
attagacctg gcgtgtcagc aaggtcatcc aggaagcaga ggctgagacg gagttaggtg
                                                                    3360
tgattactta catagtcgat tacattttac aaataacatt ttatatgtct catttactgt
                                                                    3420
gettleteec cateceatti tgtatettit eettigetti getagattig teaattitet
ctctctttct ctgtctctct ctctttcaat atctctaata atttgaaagt aattcatcat
                                                                    3480
aactaaatat ctattggggt tatgcttcac tlacaaactt ctgaaaacgg ctttactgag
                                                                    3540
                                                                    3600
atataattga tatatttaag tgtacagttt gttaaatttt gcacatattt aaaatgtgga
                                                                    3660
ctttggtaaa tgttgacata gttttacatc tgtgaaacca tcagcataat caagataata
```

3679 aacttgtcca tcaccccc <210> 2042 <211> 3641 <212> DNA <213> Homo sapiens <400> 2042 gtatgcacag tacccaggac aaatctctcc acttggaagg agatcccaat ccttctgcag 60 ccccaacatc cacctgcgca cctaggaaaa tgcccaaaag gatttcaata tccaaacaac 120 tggcttcagt gaaagctcig aggaagtgct cagatcigga aaaagctati gccaccacig 180 ctctgatttt cagaaattct tctgactctg atggtaaact tgaaaaagct attgccaaag 240 300 atetgetgea aacceaattt aggaattteg cagageeetg tgaagattea aggagaagtt ggccatctgc aaagctggaa gagtctaccc ttagtagaca ctggatctga agggcacctt 360 420 ggictiggac itcccagcci ccaaaacigi gagaigcigi itgagccaii catciaiggi gggctgttat agtagcccaa attgactatg ataaggacta aggtacaaaa tgagagttgg 480 tggagatcct gagaaagtat caggcctatt cagagatgag gaaagcttat tccaggtgaa 540 ggtagggagt ggcacaggtg agaggaatct tgggtgggtg ggtgtttatg gtaggtctcg 600 660 actaacgaat gtattcglat aatgaataag gaattgtgga agtaggagga gatgttgtat ttattctgtt tatttctaca gatctcttta ctcttttcta ccctgccttg tttccagaaa 720 780 ggctgacetg catggactgc atcaacaggc aatcttgtct ttggcttctc attgcattag gccaatgacc ttgtagatga ttagtggtgg aggaacatga acatataatg gctagatgga 840 caaaggaaag atgaatgaat aaaatcagtg gcctctgaat gttactatta ggtggcttga 900 ccttgacttt ctagtacata tttgggtaga atcatttgtt catcctctgt gatacttctc 960 cgggttttgt ttgtttgttt gtttgttttg agatggattc tcgctttgtc gcccaggctg 1020 gagtgeagtg geaceatett ggeteattge aacetacace tetegggtee aageaattet 1080 cctgcctcaa cctcccaagt agtggggacc ataggtgcac accaccacac ccagcttaat 1140 ttttgtatct ttagtggaga tggggtttca ccacattggc caggctgatc ttgaactctt 1200 tacctcaggt gatcaacctg ccttggcctc acaaagtgct gggattacag gtgtgagcca 1260 ctacacccag cctctcagat tcttatgtag ttctatggct aagttttaga agtcccattt 1320 cagggggtaa ttaatagagt catattictt ccaacaagt tgtaatctct gagctgtttg 1380 tgctcttggc acaaaagagg atgcagacag gaggatatag ttgaaaaaag aaattatgag 1440 1500 aagcattitg caaagtaaaa ttaggaggag ggaatgatga agctaaaata aatgttteet gttgaagtet gettigtatt acaaateatg aaggggettg attggatage etgetggtga 1560

caaatageet geaatleatt tetettaetg acattiggee aaaatgetge aagatacaca

1620

taaatgttac	ttgacagtgc	ctttcagcat	tttgagggag	gataaggcag	ggctctgctc	1680
aaagaaatac	ctgagttttt	ggaaccaatt	ctactgcaca	ttaccgttaa	ccctatatgc	1740
tcctttacca	atcaagggac	ctacaagata	caagtaacac	attcaaacat	gctaattgag	1800
gagacataac	aagagaacca	tctacaaagt	gctgacaggg	tttgagagaa	ccagcaaggt	1860
atgatgaagc	accctggacc	tagtatgaaa	gcaacacaga	agaaaccaga	ggtgagagag	1920
gcagaaagag	gggttcatgt	tgacgctgta	caagcacctg	gctccagtct	tgttggagtg	1980
cagcaattca	tgaagctaga	ttctccctct	acctctcaat	tatgtaagcc	agtttgtcat	2040
cttttttggc	ttgagctagt	tgaagctagt	ttttatcact	tgcaatactg	ctcatctagg	2100
ctcccttttc	cctgagtcca	tccctacagt	gctatcaatc	actttgtaca	gtgccattta	2160
ttttttgcgg	gggatgggaa	tcagactccc	ccactagact	cagagttttc	acttttcctc	2220
tttacctggg	gcctggtgca	agtttgtaag	tgtttaacaa	atacggaaag	caagcaatac	2280
aagagtcaag	gttccaagac	aaggtagttc	agtattccta	gtttcttaat	aaggtaataa	2340
ggaagatgat	gttgattatg	atgaccacca	ccactaggtg	gtagttgtgg	taatgataat	2400
ggtaatgatg	acatttacca	tttattgagg	attgcacctt	taagggcttt	acaaacattt	2460
tctcattaca	tcatcagaac	caccacctca	agtagctgtg	ttagaccatg	cttctcatca	2520
ggaagcagag	gctcagagat	ttcaggcaac	tcatccaaag	tcacacagct	agaagtggca	2580
gtcacagaat	attcactcca	aagtccatgc	tcttatccat	catgtgaata	gcccccaagc	2640
ctttctttct	acttcttcat	tttcctgaat	aaaactccct	atcctgacat	gccattcttg	2700
actctgcctt	tgcttgaact	ctatcagagc	aaggaaatag	aactaagcat	tttcctgtct	2760
cacctcctta	tgccaggcct	ggcccctgat	ataccatgtg	gcttcatgtc	aggctgagca	2820
cagaagcatc	ticacagaat	cactttgggg	cctgagaaat	atggtggcac	ctgaatcata	2880
gagttcatac	ccaaaagttt	agaaggaaca	aagcctgatt	cctacttcag	aacgtccaag	2940
ttaattcccc	aaaatatcca	atgcttcctt	agggcccaga	agcaacctaa	agcatcatcg	3000
aagcatacag	ctttgaagtc	aaatccacct	gggtcttaat	tctgactctt	tcacaatctg	3060
ggtgactttc	ggcaaattgc	atcaactggg	gaatgcctac	ctcagaaaaaa	tgatgagaga	3120
atggagagaa	ttagcactga	ccgtagtaaa	ctaatggtat	cttgcatata	gcaattattc	3180
cagcagtagt	agctatattt	attattatcg	aaatctcttg	tttttcagat	gactgaaagc	3240
caaaaaagct	tccagaggag	ttacagggaa	atgggggaaa	gataaagaat	cccgttactc	3300
cacacctcta	ctacctattg	ttccccatac	acacatgtat	atgtctccat	cttttaacag	3360
gcatgcatcc	ttctccagga	agtctttgga	ccctccttcc	cccagtggtg	ttaagagttg	3420
ccigatitac	gtaataaaaa	tatggaacac	ccagtgaaat	tcaaatttaa	ctgggeattt	3480
tatccacagt	cctagttata	cgctcctctg	cagtgtgtca	caactctcct	gtgcagtgtt	3540
tttctttctg	tattataatt	ggcctatgtc	aggagctgac	accigicaca	tctgagttaa	3600
cgtgtaactt	taagatcctc	tgatattaaa	gaattaatgt	t		3641

<210> 2043 <211> 4069 <212> DNA <213> Homo sapiens

<400> 2043

60 aaaaaggcaa gcggctctca caccctaagg tattacccag caaaaggcag cctcaggagg 120 cagcccactg aagaccitca agtccacgaa gacaatgtat ggattgttca ctaaaactga ggaatgattt tcaaataatc tgtcgccaga gggccaatcc aggcttcagg ctccagtgtg 180 tatggaggag ctgccactgc agagacgctg gcttaggggg ctgggggatg cctcctttga 240 attetgggee caccactgae aacacttete ttettggaga aaagatgaeg agaaggagag 300 gictiagaac acateettai etgaaggaca ggatacagte tigiittagg aaacteeage 360 tgctctgtgt cattgaaagg gaagaggaga gaccagatgg tccaagttcg ccatggcact 420 480 gttggtcccc tgccaaaccc agaggctata aataggatgg cagagacagt aacccatcag cacacatgaa aggagaacci gictccatca agicattitt titctatati cccigcaaca 540 atatttcgag ttcagaaacc tgtcaaagag attagttgga aaaatccctt gcctcagaag 600 aaagggaaat ctccagaaac atccagcatc ataattcatg cagcctggtg aaaaatgcgg 660 720 atacagaatt ggaggaaata gcagcatggg caccaccctg agaatgagcc taggggaacc agagagaaag cctttaccac accaagccac tctgttctca cggttctcag gatattttct 780 taagttgcca cgtccttgcc cctgtaactt tggagacttg ccctttgatc tggagagtgg 840 900 cciccigagg aggacaggat ccgcaggica gaaagaacca aiggcaigca aaiaaiggca ccaggeatea iggicaceig ccaccaegee ciccigcaac caggeeggea cigaceiige 960 1020 tglcglaalc ggalgigitc acacacgigi ggalcacata caacagigag ictaccagcc cetegeagga eegeattige iteegageit etteeeege ggagetgagg iteetaaagg 1080 ggtggagaca ggaggagctg ctgagatgaa ccatgcactc atcagccacg tggacttaac 1140 cttaaggate tgagagageg aacaacaggt ggcagccact tagaggtegg aggaggeact 1200 1260 gggggettge atggtaacat cetgaagete acaatgatgg eeegeteece attateeaca catggaaggg aaccigcaca iiiggacigi atcicicica igacgigica citiciaati 1320 ccetcatata attetitagg ggeetattet eetgaggite ticatatgta aaagggggaa 1380 aataacagta actacctcac agggttgctg tgaagaagaa acgagttgct acatagaaag 1440 caattagaaa agtgcctccc tcccagaagg tggcctgctg tcagtcatgg tggtggctac 1500 tactagacat getteacete cettgttagg ceagaagett ettgeagtee eetgggeeta 1560 ttataatatt tigcgtgcag taagtaggig gicattaaat gittiitgga igaacagagg 1620 aaacatalaa iitetigiai lalaaacati teaagitaaa talagatali tgettaiget 1680 aaaactttte tgatetttte aattataaac cacccagaaa acggttttgt gtetaaattt 1740

ttttatatca	atttgccttc	ataaattgat	accaaataag	gatctatttt	atgtcccatt	1800
aacaatggtt	ctaggctaac	tgtaaaatta	tgcaaattga	gaatttgcaa	aactgtgact	1860
agatgagggg	gcggtggaat	ggcggctctc	atctgccctg	cctctccgca	gcactttcct	1920
tttctccaca	gcttctggga	ccccacctgg	cttctctctc	accttgctac	ttctcagact	1980
catctgccca	tgggcacctc	caggagtgcc	ccaggtcctg	tcttgtcttc	atctttgcac	2040
tctccaaggt	gccttctgct	ccttgtcttt	aatacaacct	atggacacag	ggccataggt	2100
tggcacacat	ctgcctttag	ccctgactgc	tctctagaat	tgcggattct	tttctccaat	2160
gctttcttga	cactggcaca	tagacagcta	attagacttc	tcaaactgga	cattgtcaaa	2220
actctgagct	gctcaccctt	ccaagcattc	ctgtcccttc	ccccatcaa	cagcacttct	2280
gtgcttgcag	ctgatccagc	caaagatcta	ggtgtatcct	tatttccccc	ctttcctcgc	2340
tcttaatatt	cgatctatta	gcaagccttg	tcagctcttc	ctccacaaaa	taacccaaat	2400
ctgcctacct	caccccagca	cctggtttag	gccactctca	ctgtttgcct	ggatctctgc	2460
aacagtetga	tgttcctgtc	tctacttctg	cctgtactca	ctcctccaca	ctgcagccag	2520
aaatgaggcc	cactactcca	ctgcttagaa	cactctgatg	gtttcccatg	gcacttggaa	2580
taaaatgcaa	accccatctg	acttacaaaa	tcctatataa	tctggtacca	ctctgccctt	2640
tgctcagtag	gctacggctg	caagctcatt	tctgctccag	aacctttacc	ttaaccattt	2700
ccttgactgg	cctatgactc	ctgtcttccc	caacaccacc	ctctagtgag	tcactccttg	2760
tggtatttca	gatgtaggct	taaatttaaa	ctccttgaga	gaccccctga	ccaccaaagt	2820
aaccattcaa	taaccctcac	atcaccctat	ttgtttttat	ggcacctact	gttattttct	2880
tgtttccttg	tttgtctgtc	ttcctggtag	aacgtggttc	catcagagca	gggatctagt	2940
ctgttttatt	cgtcactggt	ttcacacaga	gggcattcac	caaatgtttc	tatccctgac	3000
ccactggggg	agctacagtg	agtcctgccc	caggetetee	ctgaagccta	gctggctggc	3060
tgaggagtaa	tectagetee	ctggatgatt	gctaggccat	gagacccacc	ctgagatgtg	3120
ggcatctgaa	ttaggaggag	ctggcctgca	ttctgggatc	ctgactcttg	ttacctcccc	3180
accaacactg	cccctgacc	agggccgata	gccacctgtc	gcaatgctag	aaggctgcag	3240
accagecaca	caagctttgc	tctcttcag	gctgcctgtc	ttggtgatgc	tagatgitaa	3300
acagcactca	ctgagtgctc	atgcgatgac	actgtgctaa	gcaccttcca	caagtaccig	3360
etgacecete	acagetetga	ggtggtatta	tcatccctat	tctacagatg	aggaaacgga	3420
ggctcaaacg	ggtcctggaa	gccaggtggt	ctgagaccag	agcccactct	ctctgtccct	3480
gtgccactct	gccctaaggc	ttgcttccag	ttcccagggt	actgtaaggc	tgggaaatag	3540
ggtcaaaatg	gagctgatga	gtgttaaggg	caaataatga	actctactgt	gcacactcga	3600
aagaggcttt	atatatagat	tttaactgta	aaagataatg	actaaaaaaag	tatttgggct	3660
cattttcact	tatttataca	acttgaaact	gatigittaa	atcacacacc	tctttaaaag	3720
caaaatggtt	ttaaccatca	cattttgaat	ttaaacaaac	agcaggctgc	aaacacatta	3780
gcaátcagaa	tgcgattacc	agaaaaatgc	tgttaaagtg	gaaaacactg	gaattttggc	3840
agtaatetta	gactgaaagg	gcctttctga	gtaagtcaca	gaagagtcat	ttacaagata	3900

acttetttaa ggccacaagt etgtgeteac gatgtttte teecagaata acaaagteea 3960 gtggeetaaa ttttgaaata aaaactggaa acttagatag atgttaataa agtaagteet. 4020 eetagaatea atttacetat gacacatatt taateacaga attaactgg 4069

<210> 2044

⟨211⟩ 1537

<212> DNA

<213> Homo sapiens

<400> 2044

atgetttetg agagteatgg ateteatgtg caagaaaatg aageacetgt ggttetteet 60 cctgctggtg gcggctcccg gatgggtcct gtcccagttg cagctgcagg agtcgggccc 120 gggcclggtg aagccllcgg agaccltgtc colcactgc agtglolotg gtgcclccat 180 gaccactagt gaatactact gggcctggat acgccaggcc cccgggaagg gactggaatg 240 300 galiggaaat atcittata ciggcagaac titciacaac cegicectea agagicgaci ctccctgtcc atagacacgg cgacgagcca gttctccctg agcctgcgct ctgtgaccgc 360 cgcagacacg gctatttact tctgtgcgag acatcttaat actgtcacga tttataggca 420 accetttgac cactggggcc agggageett ggtcaccgtc tecteageat eccegaccag 480 540 ccccaaggic ticccgciga gccicgacag cacccccaa gaigggaacg iggicgicgc atgeetggte cagggettet teececagga gecacteagt gtgacetgga gegaaagegg 600 660 acagaacgig accgccagaa acticccacc tagccaggai gcciccgggg accigtacac cacgagcage cagetgacee lgeeggeeae acagtgeeca gaeggcaagt cegtgacatg 720 780 ccaegtgaag cactacaega ateccageea ggatgtgaet gtgeeetgee cagtteeece acctececca igeigecace ecegacigie geigeacega eeggeeeieg aggaeeigei 840 cttaggttca gaagcgaacc tcacgtgcac actgaccggc ctgagagatg cctctggtgc 900 caccttcacc tggacgccct caagtgggaa gagcgctgtt caaggaccac ctgagcgtga 960 cctctgtggc tgctacagcg tgtccagtgt cctgcctggc tgtgcccagc catggaacca 1020 tggggagacc ttcacctgca ctgctgccca ccccgagttg aagaccccac taaccgccaa 1080 catcacaaaa teeggaaaca catteeggee egaggteeae etgetgeege egeegtegga 1140 ggagctggcc ctgaacgagc tggtgacgct gacgtgcctg gcacgtggct tcagccccaa 1200 ggalgtgctg gttcgctggc tgcaggggtc acaggagctg ccccgcgaga agtacctgac 1260 tigggcatec eggeaggage ceageeaggg caccaccace tiegetgiga ceageatact 1320 gegegtggea geegaggaet ggaagaaggg ggacaeette teetgeatgg tgggeeacga 1380 ggccctgccg ctggccttca cacagaagac catcgaccgc ttggcgggta aacccaccca 1440 tgtcaatgtg tctgttgtca tggcggaggt ggacggcacc tgctactgag ccgcccgcct 1500

. 1537

gtccccacce ctgaataaac tccatgctcc cccaage

<210> 2045 <211> 4845 <212> DNA <213> Homo sapiens

<400> 2045

60 acacaagtag gagcaataac acaaaaccca gtagagaaat atacagaagc tatcttaaat gaagtgctag tagtcccgaa catcagtgca agcaacccac aaacttcaaa ttcagcacca 120 gcaclagaig cigcagaaac gggccataca aaicaggiac aaccigagga caigciagaa 180 aciggataig teatiaegga ecaaaciegg gaigaaalga gealigaaag titeitaggt 240 agalcaaget geattgetga gatteatace gatttggace atactggata caatgaacet 300 aggaaaaacc actcagaatg gaagatcaca cttaaagaaa tggcccagat taggagaaaa 360 tgigaaaigi ttacalaici tagattigai tcagaaataa cialagiggi alcagiggci 420 480 540 ccaataccca aaaccagaga tgattatacc tggcaatctg gaactaatgc ttcagtcttt tggcaacaag gtcaaccata tcctagattc acaatcccct tcatgagcat tgcatcagca 600 tattatatgt tetaegatgg gtaegaagat gataatggta ceaectatgg ggetgetgtt 660 actaatgaca tgggaacgct ttgtgtgcgc atagtgactg agcaacagaa gaatgaggtt 720 aagataacca gtagagteta teacaagget aaacacatea gtgcatggtg tecaagacca 780 ccaagggcgg ttgcatatca acacacatat agcccaaatt ttgtgccacc aacaggagca 840 gtccaaactc acattaaatt cagacccaat gttaaagatg tgacatcagt aatgacagca 900 ggiccatcag actiglatgt acactetagt aatticatit acagaaactt gcaccigtgt 960 gaaccagaaa acttaaalga ttcagtccta attagttact ccagtgatct tgtcatttac 1020 cgcacaaata ctacaggtga tgacataatc ccaacatgtg attgtactct aggtacttac 1080 tattgcaaac ataaggacag atattatccc atcagtgtga caaaacacca gtggtatgaa 1140 atacaagaat cagattatta coctaagcat atteagtaca acatattatt gggtgtaggg 1200 ccctgcaaac caggtgattg tgggggcaag ctcctctgca aacatggtgt aattggtata 1260 ataactgctg gaggtgataa ccatgtagcc tttatagatc ttagagattt ccaagttgct 1320 gaggaacaag gaataccaga atatatteac tecctiggig aagettiigg eteiggatii 1380 gtagataaca ttaaggatca gattcaaact attaatccaa ttaataaaat atctagtaaa 1440 atagitaaat gggiaataag aattaicica gccallacca taataattag aaacaatgci 1500 gatecacata caataatage cacactaget tigitigggil geteaggite accaiggaga 1560 ttlatcaagg agaaggtttg tggatggttg caacttaatt acatacataa ggaatctgat 1620

gggtggataa	agaaattcac	agagatgtgt	aatgctgcta	gaggtcttga	gtggttaggt	1680
aataaaatat	ccaaattcat	tgattggctc	aaatctatgt	tacctcaagc	cagattaaaa	1740
gtggatttta	tcaaaaaacct	taaacaatta	ccattactag	aaaaacaagt	agatggatta	1800
agacttgcaa	cacagaaaca	acagcaggag	tatattgaca	cccttactct	aatgctagat	1860
tcatcaaata	aattcttacc	cctctatgcg	cttgaaaata	agcgaatcaa	ggaattactc	1920
aaaagaggcc	agatgatcct	tcgcacatct	aaaagaactg	aaccagttgg	tgttattttc	1980
catggtgaac	caggaacggg	aaagtcaatt	acaacatcta	tccttgctcg	aatgctcacc	2040
tcagaatcag	acatctactc	actacctcca	tcacctaaat	attttgatgg	gtatgaccaa	2100
cagagtgtag	tcatcatgga	tgatataatg	caaaatccca	gtggagaaga	catgtcttta	2160
ttctgtcaaa	tggtgtcatc	agtaccattc	ataccaccta	tggcagattt	accagacaaa	2220
gggaaaccat	tctcatcaga	ctatgtactt	gctagcacta	atcacactct	actccaccct	2280
ccaacaatta	catgcacaac	agcaatgaat	aggagatttt	tcttagattt	agacatcatt	2340
gttaaagatg	attataaatt	aggtcagggt	aaattaaatt	tgcagtgtgc	actcaagcca	2400
tgtaaggaag	ggaaaattgg	caatgcaaaa	tgttgccctc	ttatttgtgg	aaaagcctta	2460
caatttagag	atagaagtaa	tggggaacac	ttgtcccttg	ctacaatata	taataggatt	2520
acacaggaaa	gcaagaacag	aaaggaattg	acaaactcgc	tgcaggcaat	tttccaggga	2580
ccaattgata	ttgtaaacaa	gccaccacca	ccagctatag	tagatttact	taaatcagtt	2640
agaagtccag	atgtaattag	atattgtgaa	gagaacaaat	ggataattcc	agcagattgt	2700
agacttgaaa	gggatctcaa	ttatgctaat	gtaataatat	ctatgattgc	caatgtaatt	2760
agtataatgg	gtgtgatcta	cattatatac	aaattgtttt	gttctttgca	aggaccatat	2820
tcaggagaac	caaaaccagt	aacaagaaaa	ccagaaagaa	gagtggtcac	gcaaggacct	2880
caagaggaat	ttgggcgaag	ccttatgaaa	cataacacat	gtgtggtcac	aactaacaat	2940
ggaaaattca	ctggtttggg	tatctatgat	aatgtaatga	taataccaac	acacgetgat	3000
gcaggtcagg	aggtggaagt	ggatggtatt	aagaccaagg	tcagtgatgc	gtatgatcta	3060
tacaatacac	aaggtgttaa	attagaaatc	acagtactta	aactaaacag	aaatgaaaaa	3120
ttcagggaca	ttaggaaata	cattccagag	agtgaagatg	actattcaga	atgctgtttg	3180
gcactagttg	caaaccaggt	agageetaca	attttagaag	tiggigatig	ttgttcatat	3240
ggaaacatct	tattaagtgg	taatcaaact	gctaggatga	tcaagtacaa	ttaccccact	3300
aaatcgggct	tttgtggtgg	agtcttatat	aagataggat	tgatcttggg	tatacatgta	3360
ggaggtaatg	gaagagatgg	tttttccgca	atgttattaa	gatettaett	taatgaacaa	3420
caagggaaaa	tcgtatcaaa	agctgatgtg	aaagaacata	acctatatag	catacacact	3480
cctacgaaga	caaaattaca	acctagtgtc	ttccatgatg	tgttcccagg	cagtaaagag	3540
cctgctgtat	tatccacaag	agatccaagg	ttagaagtag	atttagatag	ttetattite	3600
tcaaaatata	agggtaatga	ggcagttaaa	atttcagaaa	atatgctggt	tgctgctgcg	3660
cattacacag	cccaattaac	aacactggat	attgatccac	aaccaattag	cctagaggat	3720
agtgtgtatg	gaattgaggg	tttggaggca	ttggacctcc	acactagtgc	tggatatcca	3780

```
tacacagete atggaattaa gaagaaagat ettataccaa aagacaaaaa tttaacaaaa
                                                                   3840
cttaaaattg ctatggagaa atatgggtta gatttaccaa tgataacatt tcttaaagat
                                                                    3900
                                                                    3960
gaacttagaa aaccagagaa aatcagtaca gggaaaacta gaataataga agctagtagt
ttaaatgaca cagttcagtt tagaatggca tttggtaatc ttttttctaa attccacaaa
                                                                   4020
aacccaggta ttgtcaccgg atcagcagta ggatgtgatc cagaggtgtt ttggtcaaaa
                                                                   4080
attocagtta tgctggatgg agattgcctt atggcatttg attattctaa ctatgatggc
                                                                   4140
agcctgaatc cagtgtggtt tgagcttctc gagagagttt taaatgatct cggttttcct
                                                                   4200
ggaaaattag ttaataaatt gtgccactct aagcatattt acaaaacaac atactatgaa
                                                                   4260
gtagagggtg gaatgccatc aggttgtgct ggaaccagta tatttaattc aatgattaat
                                                                   4320
aatattataa teagaacaet agttttagat acttataaat acattaatet agataagett
                                                                   4380
aaaatacttg catatggtga tgatgtattg ttctcttacc citatgattt ggacatggca
                                                                   4440
gaattageta aagaaggaaa caaatatggt etgacaatca cacetgeaga taaatcagae
                                                                   4500
aaatttgaaa aattaaatta tgaaaatgca acctttctca aacggggctt caaacaagat
                                                                   4560
gacagatata aattettaat acatecaate tatecagaaa gtgaagtttg ggaatecatt
                                                                   4620
agatggacga agagtcccag aaatatgcag gaacatgttc tttccctgtg tcacctcatg
                                                                   4680
tggcacaatg gtaaagacaa atatgattca ttcgtgaaca agattaggag tgttagtgct
                                                                   4740
ggtcgcgcac tctatattcc accatatgaa ctcttgttac acgaatggta tgaaaaattt
                                                                   4800
taaacggata tagaaagtat aaatgaagta gtttatagtt tttat
                                                                   4845
```

<211> 3764

<212> DNA

<213> Homo sapiens

agagtcagca	ggagtgagtt	caggaatcct	cgggacaagg	cactticctg	agcactggac	60
cagcgacctc	ttggcttcca	gggaggacac	acagccatca	tggaacccaa	acctcagaag	120
agtccaggta	cccgaggggt	ataatcgcag	aagcagaaat	ctttttattg	aaaatgcccc	180
acagtttcct	tcaagctaac	caggatacag	aacttggtgg	tttttgtaaa	ttccagtgta	240
gaagttggca	taagtagcca	ggaaaagatg	caatctgtgc	agaagatgit	taaatgccac	300
cctgatgagg	tcatglccat	cagaaccact	aacagggaat	acttcctcat	tggccacgac	360
agggagaaga	ttaaagactg	ggtctccttc	atgtcatcat	tccgccagga	tataaaagca	420
acacagcaga	acacagagga	ggaactctca	ttgggtaata	aaagaaccct	cttctactcc	480
agccctctcc	ttggcccttc	cagcacatca	gaggctgttg	gctccagctc	accaagaaat	540
ggtctccaag	acaagcattt	aatggaacaa	agttctccag	gatttaggca	aactcaccta	600

```
caagatttat cagaagccac tcaagatgtg aaggaagaga atcattatct tactcctcga
                                                                     660
                                                                     720
agtgttcttt tagagttgga taatatcatt gcttccagtg attctggtga atccattgaa
                                                                     780
actgatggtc cagaccaggt ctctggaaga attgagtgtc attatgagcc aatggaatcc
                                                                     840
tatittitca aagagacate ceatgagtet gtggatagea geaaagagga acceeagace
cttccagaga cccaggatgg ggacctccac ctgcaagaac aaggctcagg aattgattgg
                                                                     900
                                                                     960
tgtctttccc ctgccgatgt ggaagcacag accacaaatg accaaaaggg taatatcccc
gatgaaagcc aagtggagaa actgaacgtt tteetttete eteetgatgt catcaactat
                                                                    1020
                                                                    1080
cttgctctca cagaagccac aggacggata tgtgtgtctc agtgggaagg cccccacgt
tigggatgca tattitgcca cggagatcat citciggcag tgaatgacci gaaaccccag
                                                                    1140
agcetggagg aggtetecet gtttettace eggtecatee agaaggagaa attaaagett
                                                                    1200
                                                                    1260
accateggea ggateceaaa tteagagaea tteeatgeeg eateetgtat gtgteeetea
                                                                    1320
aaatgecaaa gtgetgeace tteteagetg gataageeta gaetgaacag ageteecaag
                                                                    1380
aggagteegg ecattaaaaa gageeageag aaaggageea gggagtaaeg cacceeagae
ccatggcage agaaccagga tggagctggg actgtccagc tctgcccct gctgctgcca
                                                                    1440
tgtgatagga gacagtcggc accccctct gaatttctgt atctgcatct taacaatggg
                                                                    1500
                                                                    1560
gatgactate ecctetetgg ttattgtate agaggatta agagggteat gtggeatgat
                                                                    1620
tggagaacct gggggaattg gaaggcctta ttatctcagc tattgtccca aacaccacag
                                                                    1680
acacagattg ggtcagtcct tcatgtaata catgctgtgt tctgtgagga tgtggtccac
acaatteett ettigitaag ggacatacag tigcaaatac teacigcaag aaggeaagat
                                                                    1740
teccaagaga gatgtgatag etgateagge tteccagaca ecteetteec aaacacetee
                                                                    1800
                                                                    1860
ttcccaacac ctccttcccc aacacctcct tccccaacac ctccttcccc aacacttcct
teccaacace tectteccaa acceptect ecceaacate etteccaaca ecteettece
                                                                    1920
                                                                    1980
aaacacctee tteecaaaca ceteetteee aaacacetee tteecaacae eteetteece
aacaccteet teecaaacae etettteeca aacaceteet teecagacae etectteeca
                                                                    2040
acaccgeett eccaacacet cetteccaaa eccettteec aaacacetee tteeccaaca
                                                                    2100
                                                                    2160
ectectice aacacetget tecceetice ceaacacete etteccaaac ateccetice
                                                                    2220
caaacacctg celetetea acceeacagg ceagagtgel gagacagagt ggceltligg
                                                                    2280
atteaataag tatettgite tettaaagae teageaaega tittagaagi egeageagit
                                                                    2340
ttacatcaca tgcagccaag atcagcttgc tctgcaagca ataacagaac tacttagcac
                                                                    2400
ticaaggiig aaagticiic actaatggat ccaligacia aligaleetg gaaggecaaa
                                                                    2460
ggaataaaat tettttatat aaataggaaa acaaaggcag agagetaaag cactaatcaa
                                                                    2520
ateggggggt gttagageaa aaacaggett cagaaagagt attitaccae getteacatg
                                                                    2580
gaaaaaateg ageeeeggag egaegaaagg catatttlet tigittetee aagitteata
                                                                    2640
accgttcagt tgcagaacca agaatctaaa accagctctg ggaaacaaat gtccagatgc
                                                                    2700
cagootcata gitgaactig gattigaaaa tacettoago actiagaaga gacaitoaaa
                                                                    2760
tacatttcat ttcctgttat ccagattgtt cggaaagtat taaaaatttt tcatttacct
```

gctgatacgg	tttggatctg	tgtccctaac	aaatcccatg	tcgagctgtg	gtccccggtg	2820
ttggagatgg	agcctggtgg	gaggcagctg	gatcgtgagg	tcatgggggt	ggagttctca	2880
cgaaggagtt	agcatcatcc	ccttggcgct	attctcgtga	gagtaagttc	tcgtgagatc	2940
tggttgttta	aaagtgtgca	gcacctctcc	gctcactctc	ttcctcctgc	tcctgccgtg	3000
taagatgcct	gctccatctg	ccgcaagtga	aagcttcctg	aggtctcccc	ggaagcagat	3060
gctgccacgc	ttcctgtaca	gcctgcagaa	ctgtggacca	atcaaacctc	ttttcttata	3120
aattacctgg	tcttggggat	ttctttattt	aatgtgagaa	cgcatgccct	tttggatcta	3180
ctgtttctac	ttttataaat	ttatcctgca	gaaatacaca	aatacacaaa	gatacatgta	3240
aaaaaagtag	tttactgcag	tactgtttgt	aataataaaa	aatcaggctg	gacgtggtgg	3300
ttcatgccta	taattccaac	cctttgggag	gccgggacag	gtggatcacc	tgaggtctga	3360
agctcgagaa	caacctgacc	aacatggaga	aaccctgtct	ctactaaaaa	tacaaaacta	3420
gctgggcatt	gtggcacatg	cctgcaatcc	cagctacttg	agaggctggg	gcaggagaat	3480
cactagaacc	gggaggcgga	agttgcagtg	agccaagatc	atgccattgc	actccagcct	3540
gggcaacaag	agggaaaccc	agtctcaaaa	aaacaaaaaa	aaaaaatcat	gtgggtattg	3600
cttaattctg	atttcatatc	attgaacact	gtagatatta	aaatgttcag	caggcacagt	3660
tctgtaaaat	tgttcgtgat	acattaagaa	tgaaagaatc	aagttgtata	ataaggataa	3720
catcatccca	cttttgtaca	aataaatgtt	tggtgtttgt	gtgt		3764

<211> 3828

<212> DNA

<213> Homo sapiens

```
aaatagagac agacttctgg caaggtagga ttatcaggga gaataattaa tgaaacctcc
                                                                     60
catgagttgg tggaaggcct atcttctaag catttcacat gctaagaagg caggtacttg
                                                                    120
tattcattit tcaaagaggg agaatgagat tcagagaagt atagtaactt gcccaaagtc
                                                                    180
ccacagetgg catteagace caaacttgag caagtecaaa geetgggtte teeegetaca
                                                                    240
gcgtgggcaa ccacagcctg cctttttaca caggctgcgc cagaggtaca tgctgtgtcc
                                                                    300
cttgagagca ctccttttac agacttattt cgtcaaaatg gcacagccag gttgcctcgg
                                                                    360
agataggaaa ccccacaatg gtaggacaaa agaaggtgcc gtgggcctaa gtaccagcat
                                                                    420
                                                                    480
caaaacaaac aggccaacca gaagtacaag gttaccttct acagcagacc ttgaaataaa
aagetteaga agggeactte tgteeettte cattaggtat aaaattteea geeetetgte
                                                                    540
                                                                    600
gtgttggggt tatttggaca gtetetegtt tteaggggta eeagtatata aaacteeaga
acgggcgcag tggctcacgc clataattcc agcactttgg gaggccaagg cgggcagatc
                                                                    660
```

acctgaggcc	gggagttcga	gatcagcgtg	accaacatag	agaaacccca	tctctactaa	720
aaatacaaaa	ttagctgagc	atggtggcac	ttgcctgtaa	tcccagctac	tcgggaggct	780
gaggcaggag	aatcgcttga	acctgggagg	cagaggttgc	agtgagccga	gactgcacca	840
ttgcgctcta	gcctgggcaa	caagagctaa	actccatctc	aaaaaacaaa	acagacaaaa	900
aacctccaat	aatacattta	tgacacgttt	tctgaatatt	tgagaattat	ttcaaccact	960
caaaacattt	taggccacgg	gcagtggctc	acacctgtaa	tcccggcact	ttgagaggct	1020
gaagcaggag	gatctcatga	gtcggggagt	tcgagaccag	cctgggcaac	gcagcgagac	1080
ctcctctcta	cagagatgaa	aaaattatcc	aggtgtggtg	gcgtgagcct	gtagtcccag	1140
ttactcagga	ggctgaggca	agaggatccc	ttgagcccag	gagttcgagg	ctgcagtgag	1200
ctaagatgat	gccattgtac	tccagcctgg	gagagagtga	ggccctatct	gtataacaaa	1260
acaaaacaga	aagacacaca	ttttaatcct	tctgaacttt	ttgagtagat	gatctgcctg	1320
gagaaataat	tctcaccaaa	ttgttaaaag	gttatgaaag	ggaatttaac	tcagttattc	1380
ttaatcatga	tactctttat	ttttagttcc	ccatttgtat	tatgttggga	ttttgatgta	1440
attatcacat	cacttgcatt	gatctttata	ctctccatgt	acttgaaaaa	gaaatagcaa	1500
catattttta	agggctgggg	cacccagcat	tcaaatgaaa	atccaggatg	aaggaagaac	1560
aaaagatcat	ttcattgtcc	ttccaacacc	agctcagagt	gaaagctggt	tgagttaaat	1620
tccttgtgaa	atgcattaat	gacagtagca	gattttactg	agcatttact	acattcccag	1680
cactgtgcta	aatgtgtcgc	aagcatgctc	tcacttcatt	ctacaaaatg	aattctcatt	1740
ttccagatga	agaaactgag	gcatgagaca	taaagttagg	tagtatgtcc	aaagtcatgt	1800
ggtctctatg	ctattgaacc	agaatttgaa	tcctgctggt	ttcactctcc	ttgccaacca	1860
ctaccccaag	cacatecege	ccctactgtg	tctcgtactt	gctcttctct	ctgcctgcag	1920
cacctctgtc	tggttttctc	cagccagctc	cttctcactg	ttcaggtccc	aaccaaaagg	1980
cacttcctta	gggaggcttt	ccctgaccat	cctacccatt	gtgtccccag	ctccaccaca	2040
cagcctctgt	catagcaccc	atcactgcac	ttgagcacca	caggagacta	tttactcacc	2100
tgtcctttgg	ctgcctcgcc	tgctataata	tcagagccac	aaaaacaggg	ccttgtatct	2160
attattcacc	actttatccc	cagggctcaa	cacagtgcct	agtacatagt	acatgctcag	2220
taaagttgtg	atgattgagg	gaaccctgcc	tccactgtat	acagtgcaga	acaccaagcc	2280
agggccagga	aaacccctga	cggtccctag	gtctgagctg	ggagcaagag	gaaagggaat	2340
gaacagtaac	cctttgatgt	attcagtaac	tgtctaatga	gtcccttgtg	ctaagacttc	2400
taggggatac	caaaaacatg	tccctttctt	tctaagattt	aaagagtatt	tgaggaggtg	2460
aaaccatcat	ggtaaacatt	gtcgtacccc	tcaaaacatg	cccaaatgtc	aaaatatggt	2520
atgcaattca	gatgctaaac	tgataaaaga	gacagcactt	gtattaatag	cattgtcaaa	2580
atgcactggg	gataaaatac	agaagaagag	tccacacact	gtttcacgag	aaggagtgta	2640
tcatgatttg	tagtaatcga	agaacatgtt	tatgggaaca	gggtgactca	gctctcctgg	2700
ggaggatgga	tgaggagtta	gcaggaagag	agggtaccaa	gtgaggggaa	agcagcaggg	2760

```
tgggtctggg gcatggacag gaaacagagg ctgggaaaag ctacatcttt tattcatgct
                                                                   2820
ttttcacagg agctgaagtg ggaatcagta catcgagaat ccacgcccgg ggaccagtag
                                                                   2880
gacttgaggg actgcttact actaagtggc tgctgcgagg gaaggaccac gtggtctcag
                                                                   2940
attictcaga gcatggaagt ttaaaatatc ttcatgagaa cctccctatt cctcagagaa
                                                                   3000
acaccaactg aaaagagcca ggaaaacccg ggaattttcc aaaaggtctt cacgttaaac
                                                                   3060
tigicitate teaggagaga geeegeteet gieteeagt teetggtagg gietgeetgt
                                                                   3120
lggaaagtgt acctggatgc ttctgggctc cgtttggcaa tagcaatctt ggctgatgtg
                                                                   3180
cacagtotgg otoccagoto accotttttt tttaaagtaa gaaaatagtt gotacogata
                                                                   3240
gggactttgc caagtccaat tatettetag gattgaaagg tgcattttcc ccataaaaaa
                                                                   3300
ggcgaggaaa acccatggct gctttgtgtc acctcagtga cttacagtcc cccttggcat
                                                                   3360
ttagttggta ctagagccag tcatccttaa caaatctttt cacattttat ttctttcaca
                                                                   3420
tgcagtcatc ttcaaaaagg aaagatttgg aattttagaa aaggggcaac tcttctttt
                                                                   3480
agcattetea teagaaagte acaaaaateg atggaateat ttecaetggg aagattgace
                                                                   3540
ttttgtattt atttgtgggg taaattaata agcattccag atgcttgcag cttcctgcat
                                                                   3600
ccaggagatg ctgtgttccc cgtgatgcag ctggaaccca agctgcagca ggagatgcaa
                                                                   3660
gilicaggai gilccccaci gagciggagg aataictaca gcagigaigi iigaaattii
                                                                   3720
tgtatgaatt attttgtcgt cctacccttt tcctccaaaa caaaaattag aggattattt
                                                                   3780
taatacttig gattcticcc cctttttiga gaaataaagt tttttatg
                                                                   3828
```

<211> 3894

<212> DNA

<213> Homo sapiens

ctcatcctgg	ctgctctcac	cgtggcctgt	ccagatgcag	gagctcctct	ctgaatctgg	60
ggctactggc	agaaccagta	aacacggagt	tactcctgta	ctgagctgag	taaaataatc	120
tgactgagag	gatgcgctga	cctcagtttc	gacaactgcg	tttggtacca	agccctgcaa	180
gggctccacg	gagcagcttt	gggggagacc	tgcctgcagg	aacatgtacc	ccacggagca	240
gctttggggg	agacctgcct	gcaggaacat	gtaccccacg	gagcagcttt	gggggagacc	300
tgcctgcagg	aacatgtacc	ccacggagca	gctttggggg	agacctgcct	gcaggaacat	360
gtaccccacg	gagcagcttt	gggggagacc	tgcctgcagg	aacatgtacc	tcacggagca	420
gctttggggg	agacctgcct	gcaggaacat	gtaccccacg	gagcagcttt	gggggagacc	480
tgcctgcagg	aacatgtacc	ccacccgaca	cgtcctggga	gcctcgtctg	aggtacaaac	540
aacaggaaag	cactgatgca	tttttcaaaa	tccagcagga	gggaacggtg	ggctgtggat	600

gctggctggg	aaagctcctc	gggcacagcc	ctgtgggcag	ggaggggagg	agggctcagc	660
ccccacacag	gccgcctggc	accaggagtc	acaggcctca	gccgtgggat	gtccccagag	720
ttccaaccgc	cactcttgca	gaagcagccc	agcagggtga	gggtggggcc	acatggggct	780
cagctgcagg	agggacgcca	ggtcctgcac	ttctcacccg	cagtgacctt	gggcagggca	840
ttcattcctt	gggagaaatt	tcctcgttgg	tgaaatgaaa	tcactgcttg	gcttcagcca	900
cataatgtta	ggcacgctaa	ctgcagccta	ggcaacctca	gaccctcagg	aaatcaacag	960
aggggtgcca	gctccctgca	caggtcccgg	cctaactcgg	gatgccactc	agggccctcg	1020
tcttcccatc	ctgtggctct	gtcttcacaa	ggccccagag	gtgctcttgt	cccttccctt	1080
tcagtccctc	agccagtggg	cagcacacgg	ccacccaaac	acaagaggcc	aggaccatgg	1140
acagcaggga	gcacagagcc	caggcctccg	tgatcctagg	aacacgcagc	atccgggaac	1200
acggaaagta	aagatggaga	catggggcgg	gaggaagcta	agcagggaca	cagtaccccc	1260
ttgcatcacg	gaaatgcctg	gccagagcga	cctgccgcaa	gaagccagcc	cagctgctcc	1320
tgtccctgaa	atgtccggag	agagggctag	cagggaggct	ggcgcctggg	ccaagagagg	1380
ggctactcag	ttcttccaga	acattccagt	gtggcccatg	gacaccggcc	ttctgatgtc	1440
cagagagggg	ctactcagtt	cctccagaac	attccagtgt	ggcccatgga	cgccggcctt	1500
ctggggtcca	ttctgtcctg	tgtcacttca	gttgatgagc	tgcttgagac	cagaactgcc	1560
caaatccaga	accgcccact	accttctgtg	aggctgtggc	cagaaagcaa	gccagacttc	1620
tgaagctgcc	tgggcctgtc	gggacccagg	agaatctggc	cgtgaaggag	aataaaggag	1680
gaagccaggc	ctggcacagg	gacagggtgg	ggacccagtg	agatetecaa	ggaggaagcc	1740
agggctccta	cactggggct	gctgttctcc	cggaggaact	ccacccaagg	agagtctggg	1800
attatcatga	gagacaggac	cgcatctgtg	cacagtgcag	tacgtcaggt	gctggccagg	1860
ggccgggggc	ctcagggagg	agagtcaccc	accaggccaa	ctaggacaga	cgaaacgtga	1920
gtgcccctac	gggagaaagc	aaagctgaga	cagcatcgcg	agctgaggga	gaaactgaca	1980
gacggcagtt	caccaaaaacc	caaaaactgg	tcattctctg	gcttttaaca	aaccaaagta	2040
tatttctccc	tctgaaataa	gaaacacagg	acaattatta	agttccaaaa	gtacgtttca	2100
ttttggaggc	atgttgttgg	tcccctttgg	aatcatgaac	ccctgtgagc	gaaacacctc	2160
ccaccattga	ttctgacagg	gtacggcggg	cagttcccgg	cccaggtaga	ggcagacagg	2220
tgcagagcca	cagggccacc	actgcagagt	ctggccttct	ctccagcccc	gggtgcaccc	2280
acggttatca	gggacccagc	actgcctccc	tgcacgcaca	tggctctcca	ggccaccact	2340
gcagagtccg	gccttctctc	cagccccggg	tgcacccacg	gtgatcaggg	acccggtgct	2400
gcctccctgc	acccacccgg	ctctccacag	cagcaaacgg	ggtacattag	ggtggacggg	2460
atgtggggcc	agggccctgc	tagggctggg	gtggactgcg	gagggccggc	accaagcagt	2520
tccaggtgtg	gagggcggcc	ctatgtcagc	tgttagacac	gcaggggagg	cacctcagat	2580
ggctacaggt	ttgattgtgt	ccccacaaaa	atccatatgt	tgaagtccta	acccccaaca	2640
ctgccgaaga	tgaccttatt	tggaaataga	gtcatcaaag	acatcattgg	ctacattaag	2700
atagggttat	actagagtag	ggggacacct	agcttattat	gactggtgtc	cttataaaaa	2760

gaaggaaact	ggacacataa	agggagaatg	ccataggagg	acggaggcgg	agatcggggt	2820
gaagcttctc	taagccacgg	agagcggcct	agaaccgacc	cttccctcac	agccctcaga	2880
ggacagcctg	gaaccgaccc	ttccctcaca	gccctcggag	gacggcctgg	aatccactct	2940
tccctcacag	ccctcggagg	gcagcctgga	accgaccctt	ccctcacagc	cctcggaggg	3000
cggcctggaa	ccgacccttc	cctcacagcc	ctcggagtgc	gacctggaac	caacccttcc	3060
ctcacagete	ttggagggaa	cccaccctgc	ccacaccttg	acctcggaca	ggtggcctct	3120
agagacctgt	gcagtgagtt	cctgctccca	gcctgtggtc	cttccatgtg	gaagcaaagc	3180
aaactcctcc	aggcacattc	accgccattg	gcatgggcct	ccgacactga	ccagggcctc	3240
ccgtcacctc	tgcccctgcc	caccactccc	cagcccaggt	accatgctgt	aaaaacagcc	3300
tcaaaaagaa	catgaggtcc	acagctcctc	caggagactg	ggccagcccc	aagcacatcc	3360
agagaggtgg	ctcctctgac	tggaggctca	cgccaaagcc	acacagagac	agctgccatt	3420
ctcgctcgct	catgcttccc	ccgagcctaa	accctgacca	gccagctcta	tacatttaca	3480
tctttttctg	gcctcacaca	ctgtctagaa	tgtccagtcg	aatgttgaga	agtcgtggtc	3540
aaagcagaaa	gcccagcttt	atccccagtc	ttagtgggta	cgtgtttgct	gtttcacgtt	3600
aagatactgg	ctggcagtgg	ggcacagtga	ctcacgcctg	taatcccagc	actttgggag	3660
gccaaggtgg	gtggatcaca	aggtcaaagg	attgagaccg	tcctggccaa	catggtgaaa	3720
ccccatctct	actaaaaata	cagaaattag	ctgcgtgtgg	tggcggacac	ctgtagtccc	3780
agctactcgg	gaggctgaga	ccggagaatc	gcttgaacgt	gggagcagag	gttgcagtga	3840
gccgagatcg	caccattgca	ctccagcctg	ggtgacagaa	cgagactcta	tctc	3894

<211> 4331

<212> DNA

<213> Homo sapiens

aagaattgat	ctacccacaa	tgtcaacaag	tacccctttg	aaaaacgcta	ccaactaaat	60
gggctttggc	aggccttcct	gagaatctaa	acacaatttt	taatgtggtt	gctctggcag	120
agactgctgt	ctcatcagcc	tatttttaga	ctaccaaaca	agtatgtttg	aattataaat	180
ttaacctcca	cacccatttt	tcttttttta	actttttatt	atggagactt	ttctttttt	240
tttgagatgg	actcttactc	tgtcgcccag	gctggagtgc	agtggcagga	tctcagctca	300
ctgcaacctc	cacctcccgg	gttcaaccaa	tcctccctgc	ctcagcctcc	tgagtagctg	360
ggattacagg	tgcccaccat	cacgcccggc	tgattttgta	ttttttagta	gagatgaggt	420
ttcgccattt	ggccaggctg	gtcttgaact	cctgacctca	ggtgatccac	ccacctcgac	480
ctcccaaagt	gttgggattg	caggcgtgag	ccaccatgcc	tggctgagac	tttcaaattt	540

```
atataaaagg gagaaattag ccacccagcc tcaacaggtt ttatcaattc tgtttcatta
                                                                     600
tetecateae caccaacace tettegtett etaattgetg gagtatttta atgtaaatet
                                                                     660
                                                                     720
calcetatee titeaaceaa aattietgea alagtgaeta alacatgeee titittitga
aacatcatta tacgtaacag ttgacagcag ctcttaagtg tcatctaata tcctatttca
                                                                     780
                                                                     840
tgtacagatt tatcagattg acccagaatg tetttttata gtttttttgc tttgtttgt
                                                                     900
tttacagtgg tttgttcaaa catggattca gataaggtcc acacatttta gtctgtaata
                                                                     960
gtttcttctc accetetete accettgttt teettetatg teatttattt gttgaagaaa
                                                                    1020
ctggatcatt tttcctgttg tggaattcca tattctgggt ttggctgatt atatgtttct
ctgtctctct tactttccat gaactggtgg ttagacataa agactttcag aactgattgg
                                                                    1080
taagatatac atttatttcc attggattgg aagtcataat atctgattat cccctttttt
                                                                    1140
tttttttggt catgttgaga ttgattatag tagttcagct gttgtaagtc tattccaccc
                                                                    1200
ataaagttcc tcagcaaact ttaacctaat ggttttaata gtcattgatg atgtttaaat
                                                                    1260
ccatttcatt aaatgctgca aaatggtgat attctaattt tttaaattct aacttctgca
                                                                    1320
                                                                    1380
ttcgttagct ggagttttt ctacaaagag ggactttgcc atatcagcta tttgcttcaa
ttgtaatatg taatgaaaag gcaggattag gtgcttgttt actcatttgc agaataataa
                                                                    1440
catteetiga aagigaccag iggggittia gggittiigt ittigtiiget iteititeat
                                                                    1500
ttigitttat taigagatca iggittiigi tgiggitgit gitatigiig tigittigia
                                                                    1560
ttggttatat tttagtccac tcagtccact aatatcactt agtttttatt acggaaaatt
                                                                    1620
                                                                    1680
tcaaacactc tcaagtagac agagttgcac catacagtga aacctcttat gttcattctc
taacgtcaac agtgatctla acattcaacc aatcttatct tcatctatac ctgtactcca
                                                                    1740
gccccacttt cttctgccct tattttagtt tgatgcatat ccaatcagtg ttcaaattta
                                                                    1800
aaatggteta aaatatttta aaaateagat tgettgaate aaaatteaga tetaceaett
                                                                    1860
agtacagttt atattgtgat atgtccttga gtataatcta tggacacccc ctcaactctt
                                                                    1920
gcaatttatt taagtaagtt gaaacattta gtcactagag atttccacgt actagatttt
                                                                    1980
gctgatttca titaltiggi alagittaat gtattttctg taaattggta gagtcaaaaa
                                                                    2040
gaaatagagc gtgggcctag ttggaaagac agatttcatt cagtactatt gcaatagggg
                                                                    2100
aaaatagaac caagtteeat tteagaatac aacaaagaca ettggggatg aageagagtg
                                                                    2160
                                                                    2220
agagggtcaa tggatggaaa ctttctaaaa ggagacatca aaggtagaag gtttctttct
gaccigacti aggaticcig ciaaaggcag gccaaggiga icaiagatcc agagtgggag
                                                                    2280
atagtttagg aggattetta etatatataa etgagetaaa eagaetgatg aeggggetea
                                                                    2340
aggacaaata clagiigatt gcicagagca gccigcitaa aagtaiggic aaggagagaa
                                                                    2400
tetttagtgt agaatggtga teagatttaa gtttgttgte etttggttet tgttttettt
                                                                    2460
etgaaaagca agacetgett caaaggtggt ggtgtgetet ettgeactag gaggtatatt
                                                                    2520
                                                                    2580
atgicitgia licaggetai tigcattica gattacacag tittatgiaa cigcittaac
tttgtgtttg tactgaatat tagtttcttg atggcagaga acatatttca ctttcagaat
                                                                    2640
gtttttctgc ttacatggat ttattttcaa gaaatttcat acaatacttt atttagaaga
                                                                   2700
```

```
2760
aagcagaatt ttetgaaate acagtatgca gaggcattta ccatcaacte tgacaaacat
ccttctggtc ccttttctat gcatgtattc tgtggaattg gatgcaaaca catattaaaa
                                                                 2820
                                                                 2880
atatatacat ttgcctaatg gaaccacagc atacagagta ttttatagtc tgcttttcca
ticagigata ticcaggaaa alattiicti atcagigigi tiagatacac alcetticaa
                                                                 2940
                                                                 3000
taggicatea ittaaattie taetgietaa eattattita aaagtaagti tiictetaat
                                                                 3060
aatcagcacc acattaaaca tactgtgtag ctttcacttt aaaattattt ttatggacat
                                                                 3120
ttgatateat tagettgaea ttattaataa eagttaeett gaetttttga tateatetgt
actgtcttgg aaagtgaaaa tatttgtcaa actgttaaat gataagaaag aataattata
                                                                 3180
                                                                 3240
cactgccaag cagaatttcc ttcttttgct ccctcccac cttctgctcc aatcacataa
ataagagctg ttttttcttt gcagtatgca ttgcctcagg aacaaaggtg gctctgttta
                                                                 3300
atcgactacg atcccagaca gttagtacca gatacttgca tgtagaagga ggtaattttc
                                                                 3360
atgccagtic acagcagtgg ggagcctttt ttattcatct cttggatgat gatgaatcag
                                                                 3420
aaggagaaga attcacagtc cgagatggct acatccatta tggacaaaca gtcaaacttg
                                                                 3480
                                                                 3540
tgtgctcagt tactggcatg gcactcccaa gattgataat taggaaagtt gataagcaga
ccgcattatt ggatgcagat gatcctgtgt cacaactcca taaatgtgca ttttacctta
                                                                 3600
                                                                 3660
aggatacaga aagaatgtat tigtgcctti ctcaagaaag aataattcaa titcaggcca
ctccatgtcc aaaagaacca aataaagaga tgataaatga tggcgcttcc tggacaatca
                                                                 3720
                                                                 3780
ttagcacaga taagttgaat ggcggtgggg acgtagcaat gcttgaactt acaggacaga
atttcactcc aaatttacga gtgtggtttg gggatgtaga agctgaaact atgtacaggt
                                                                 3840
giggagagag taigcictgt gicgicccag acattictgc attccgagaa ggitggagat
                                                                 3900
gggtccggca accagtccag gttccagtaa ctttggtccg aaatgatgga atcatttatt
                                                                 3960
4020
caggagcaat cettegagee aatteaagee aggtgeeece taacgaatea aacacaaaca
                                                                 4080
                                                                 4140
gegagggaag ttacacaaac gecagcacaa attcaaccag tgtcacatca tctacagcca
cagiggtate claactaceg tettitiget aggacitaaa etgaciigag igiggeaaaa
                                                                 4200
agitaacaaa aaaggagaaa aaatgaacaa tegiitgigg titetiggga aaaciittea
                                                                 4260
laccaggiga tactaticaa aaaccccgti gictccctgc aagigcigat tigaaaigca
                                                                 4320
gaagccacag t
                                                                 4331
```

<211> 2538

<212> DNA

<213> Homo sapiens

```
60
tttttaggag cacgggtact acttactgtg gacgacggtt ggtcaaggaa ggctttctgg
                                                                     120
aggaggtgac agctaggctg ggtcttaagg atgaatggga agagaggga gaacatgtgg
                                                                     180
ataaggccag gcaaaagggc tgcacagcca agtcacagcc aagacgaaat gcagggagag
                                                                     240
ltctggaagc tgcgtgtttc atgctgctgg gtagtgtgga aggacaggct ggagctaggc
agctaagcag cttggcaaat ggagctactg aggattccaa acaggacctc tgcagtcgtc
                                                                     300
                                                                     360
tccactgctt atgggttgaa ccacgtgaaa tagacaatat tcggccattt agggccaaga
                                                                     420
caaatgccag ctttgcgggg tgcagcctca cagagaggct gcttgggggc ctttgcagag
                                                                     480
ggtggatgag cagagggcat cctccggaac ctgcttgggg acccggctct gaggccatcg
ggccggtggt gtccagattc tcgtgtaggc tgggagaaag gggaggttca agaaacacgg
                                                                     540
aggaagtgaa gcgtcagagc cggggggacg gggtgccgca gaggagaagg agcactgagg
                                                                     600
                                                                     660
ctgaggtcca ggcttgcaga cacgtggacc atgagtattc tgccaggtct gtgggtgtct
                                                                     720
ctlctgagct acaccagttt ccaggttacc tgggaccatg gataactctc agatcagcaa
                                                                     780
ctigicagit gailliceaag ctgcigtigg ciggacteag acteageagg gagcaceigg
gegagecetg tgetgeggge tggacteegg cecatetege tgattactet tgettttget
                                                                     840
                                                                     900
cccagtgtg tcctcaagag glcagagcct gcttgttgtt tcttcatgac cacgggagga
                                                                     960
ggggcaccaa catgagggtg ctagcatctc cccagtggtg gcttcccagg gctggggaaa
                                                                    1020
ccctggggga ggggttggga cagggacctc tgtcgcttgc tgccactgcc tgggtcaact
                                                                    1080
geetggeagg getggeeget egtgeteaga aggetgagge ettacetgee tteteetet
acceagegce catglaagga cacatetgag ttggcattet gtgtetgete ttgagetact
                                                                    1140
                                                                    1200
cgcatgataa gtctttgttg tcctgtggga tgtcaccggt tcatgctgaa gagaaattgt
                                                                    1260
aaaggactee titgeetget eaggeeecat ggeetetgte atgittigte eeegteeett
                                                                    1320
tgggagcaca gcagcagtgg gctggctgga ctgtgcaggc gaggttcaag gatgaggtac
                                                                    1380
agttgtgtga aaggtgagcc tgctggaccg gggagctttc ctcaaggcct ccgcctggct
atgatggcgt tagggttgag gggaagcttc atccaaaatg cacagtactt ggatgtcaag
                                                                    1440
atgatgttgc tgctctcagg atgagtcact ctccaccact gacttccttt gatgttctga
                                                                    1500
gctcagcctg gagtctgacc tgggactata gcacttgttc tcccaaggta aggctggcgg
                                                                    1560
                                                                    1620
ccaaacccag ctgcgcacac ctgaacctgc tccttggcag agatgaaggg cgtcatgttt
                                                                    1680
cgtagccact caacacccat ggacaatttg gctccttgta aagacttagt catgcctttg
                                                                    1740
aactgactta ciigaaatat aatigcicci attiigcicc aaagaccagi ggcaigaigg
                                                                    1800
gilagagila liligialila ligagaligi tglaatlagc aalcicaggg cicagictaa
                                                                    1860
ctgcattatc catgctggaa aacttaaaaa aaaaatacag tccttcatct tcagttttcc
                                                                    1920
aalggtegee agitalaeae agetaatett tgeagtgaaa gitgtettig gagaatgige
                                                                    1980
tttcttggtc ccgggtggtc ctggtcttgg gctggaatct acgtgagctg ctttgaagta
                                                                    2040
agetgacaat acacaattat taaggetatt tigacetgea agtatggtit ettaaaaagg
                                                                    2100
aacaattaaa taccatgtag cagttattta gactttagca ttgactaagg aaaggagaaa
atggaagaag aaccccctcc tgcttagatg cagtcatttt tttaaaaagt aatcttttgg
                                                                    2160
```

ggaataaact	taaccaagga	ggtgagggac	ttgtaaacaa	aatgttaaaa	ctgcactgaa	2220
gactagaaaa	tgttgatgaa	agctgttaaa	gaagacacaa	ttagatgatg	aaaacacatc	2280
ccatgttcat	ggattgaaag	acaatattgt	taagatgtca	atactataga	ttctatgcaa	2340
tccctgtcaa	aacccaattt	tttttcaaac	ataggaaaat	ccattctaaa	atttacatgg	2400
actctcaagg	aaccctgagt	agacaaaaca	atcttgtaaa	agaacaatgt	tggagggctc	2460
acactttctg	gtttcaaaac	tacagtaatt	aaaaagctac	agtaattaaa	acagcatgat	2520
attgtcacaa	agatatag					2538

<211> 1766

<212> DNA

<213> Homo sapiens

<400> 2051

60 ageteteaga eaggtgtett ageeetggat teeaaggeat eteetetegg tgateagete tgaacacaga ggactcacca tggacttggg gctatactgg gttttccttg tcgctatttt 120 180 agaaggigtc gagigtgaag igcaacigga gcagicgggg ggaggccigg taaagccigg 240 agggtccctg agacteteet gtgeagecte tggattetea eteagteett atgaagtgaa 300 cigggicege egggeteeag ggaagggeet agagtggatt geetatatta gtagtagtgg 360 gagtaaaaga tactacggcg attcagtgac gggccgcgtc agcatttcga gagacagcgc ccagaactca gtctctctgc aaatgagtgg cctgagagtc gaggacacgg gtgtttatta 420 480 tigigogaga giogacigga atcacticia cittiticatg gaigiciggg gcaaagggac caeggicate gieleegeag ettecaecaa gggeeeateg gielleece iggegeeetg 540 ctccaggagc acctctgggg gcacagcggc cctgggctgc ctggtcaagg actacttccc 600 660 cgaaccggtg acggtgteat ggaactcagg cgccetgacc agcggcgtgc acaccttccc ggelgiceta cagiceleag gaetetaete ecteageage giggigaeeg igeeeteeag 720 780 cagettgggc acceagacet acacetgeaa egtgaateac aageecagea acaceaaggt 840 ggacaagaga gitgagcica aaaccccact tggtgacaca actcacacat gcccacggtg 900 eccagageee aaatetigig acacacetee eeegigeeca eggigeecag ageeeaaate tigigacaca cetececcat geecaeggtg eccagageec aaatetigtg acacacetee 960 1020 cccglgccca aggtgcccag cacclgaact cclgggagga ccglcagtct tcctcttccc eccanance anggatace tratgatite eeggaceet gaggleaegt gegtggtggt 1080 1140 ggacgtgage cacgaagace eegaggteea gtteaagtgg taegtggaeg gegtggaggt 1200 gealaatgee aagacaaage egegggagga geagtacaac ageaegttee gtgtggteag egiceteace giceigeace aggaciggei gaacggeaag gagiacaagi geaaggicie 1260

Ĺ

1320 caacaaagcc ctcccagccc ccatcgagaa aaccatctcc aaaaccaaag gacagccccg 1380 agaaccacag gtgtacaccc tgcccccatc ccgggaggag atgaccaaga accaggtcat cctgacctgc ctggtcaaag gcttctaccc cagcgacatc gccgtggagt gggagagcag 1440 1500 cgggcagccg gagaacaact acaacaccac gcctcccatg ctggactccg acggctcctt cttcctctac agcaagctca ccgtggacaa gagcaggtgg cagcagggga acatcttctc 1560 1620 atgctccgtg atgcatgagg ctctgcacaa ccgcttcacg cagaagagcc tctccctgtc teegggtaaa tgagtgegae ggeeggeaag eeceegetee eegggetete ggggtegege 1680 gaggatgctt ggcacgtacc ccgtgtacal acttcccggg cacccagcat ggaaataaag 1740 1766 cacccagege tgccctgggc ccctge

<210> 2052

<211> 1727

<212> DNA

<213> Homo sapiens

<400> 2052

60 atagggtagg ggaggccctg ggaaaggcag gacctcgagg cgcggccgcg cgaggtgacc 120 ggagtcacag ttcccgcagg cggcgacagc agagcgccca ctgcctccag cagattaata 180 ttaagattgg aagtttgtgt cittitgctgg atattggaaa itgaatgtaa iggcaacaga 240 attlataaag agitgcigtg gaggaigtti ctatggtgag acagaaaaac acaactiitc 300 tgtggaaaga galtttaaag cagcagtooc aaatagtoaa aatgotacta tototgtaco tecatigaci leigilielg iaaageeica geitggeigt aeigagggit attigetite 360 caaattacca totgatggca aagaagtacc atttgtggtg cccaagttta agttatotta 420 cattcaaccc aggacacaag aaactccttc acatctggaa gaacttgaag gatctgccag 480 agcatettit ggagategaa aggtagaact ticcagitea teccageacg gacetageta 540 600 tgatgtgtat aacccattet atatgtatea geacatttea eetgatttga gtegaegett 660 tecteccegt teagaagiga egagacigia iggaleggit igigalitaa ggaegaacaa acticccggi icccciggge taagcaaate taigtitgat citacaaact catcicageg 720 780 atteatecag agacatgatt cattgtecag tgtacccagt agttettett caaggaaaaa ticicagggg agtaacagaa geetggatac aattacteta teaggagatg aaagggactt 840 900 tgggagacig aaigigaaat igililataa ticticagia gaacagaici ggalcacagi 960 tttacagtge agagatilaa gitggeeete tagitatgga gacacteeta etgittetat aaaaggaata citacaligo ccaaaccagi gcatticaaa tottoagooa aggaaggito 1020 1080 caacgitige caigeagaac tegaaligg gaetigtitt caageagiaa alageagaat

tcagttacaa	${\tt attcttgagg}$	cacggtacct	tccaagctca	tcaacacctc	tgactttgag	1140
ttttttcgtg	aaggtgggaa	tgtttagctc	gggagagttg	atttataaga	aaaagacacg	1200
cttactgaag	gcctccaatg	gaagagtcaa	gtggggagag	actatgattt	ttccacttat	1260
acagagtgaa	aaagaaattg	tttttctcat	taagctttac	agtcgaagct	ctgtaagaag	1320
aaaacacttt	gtgggccaga	tttggataag	tgaagacagt	aataacattg	aagcagtgaa	1380
ccagtggaaa	gagacagtaa	taaatccaga	aaaggttgtt	atcaggtggc	acaaattaaa	1440
tccatcttga	agacticaca	cattaatttg	gtgaagaact	tgacattctt	ttagaagact	1500
tatgatttca	atttgctacc	aatgagaaga	ggcaaatcaa	caaatttgtc	aatttatggg	1560
ggctataatt	atggtatata	atgtatctga	tagaaaattt	gataagaaaa	tgtaatgaat	1620
tttatcagat	atccaaagta	aaggaaatgt	tttaaaactg	caacaagaga	cacagacagt	1680
aaaatcaaag	tattattagg	atgactaaat	aaattataaa	gtctgtg		1727

<211> 2079

<212> DNA

<213> Homo sapiens

```
cagtitggca teactectee cacaatttaa aaacecaaaa ecaacacete gtgaagetat
                                                                     60
caeggeeeag agettaaaaa ettaaaeeag gaetaaagge aceaeetgii iteaatgeag
                                                                    120
cgttgcccac aggaatcact ctgacaaccc tcacttttct aacagacccc tggcgggcag
                                                                    180
aggactaatt etettitte acattettie tgigttitte acagatgaga gagaggage
                                                                    240
tectgaggag geteaaggea ggegetgaga ggaggeaggt cegeageeag ggeeeetgea
                                                                    300
gccacagggt tccgtgcaca gcatttttt acactcaaag gcttttttat gtctttctcc
                                                                    360
taaaltgtgg taaaatacac taacaltcac citcclagec atalttaggi gcacacaagg
                                                                    420
geacaggaag tgeatecaea etgigeaget getgeeacea ceaceatete cagaacgite
                                                                    480
teatetteec aaaeggaact etgteeceat taaaeaceaa tteeceatee eeetggeeta
                                                                    540
ggecetggea teccecaget aegttetgte tetaegaagt caetgeteta gggacegeat
                                                                    600
gagtggagcc acacaggatt tgtccaggtg tctggcccgt gtcactgagc accatgtcct
                                                                    660
                                                                    720
caaggigcat gigtgctgct tiatgcatca gaatticatt cctitctgcc gittgatggc
tgaataatat tecactgegt egacagacca cattlegtti aattaggeat ceacceatga
                                                                    780
acatetggge tgtttetaac titeggigat tgtggatagi getgeeatig gacatgggig
                                                                    840
gacaggiaco totilaagao coagciiica atiototggg giotgiacoo agacgiggaa
                                                                    900
elgelgggte acagagtaat tecatettet tilgtgillt gaggaactte ceacagtgee
                                                                    960
```

```
cgcactactg tacattecca ecageggegt acaaggetee aacgteacca egecetgeag
                                                                   1020
acactetttt teetttttgg ttatttatge atacataaat aatgatgtat geattattta
                                                                   1080
tgaatgaatg aatgaacgac agggtetege tetgttgeec aggetgeagt geagtggeaa
                                                                   1140
gateteaget caetgeagee teaaacaeet gggeteaage gateeteeca cetttgeete
                                                                   1200
ccaagtaget gggaccacag gtgtgcacca gcacgtctac ctaatttttg tatttttgt
                                                                   1260
agagatgggg teleacaatg ttgtgcagge tggteteaaa cacetggget caagtgacee
                                                                   1320
teccaecteg geeteccaaa gtgetggaat tataggeeta agteaccagg ecaccaggee
                                                                   1380
agtctgttta tttatttatt tacagagtct cactctgttg cccaggctgt agtgcagtgg
                                                                  1440
catgatettg geteactgea accteegeet eccaggitea agigatiete eigeeteage
                                                                   1500
ctcccaagta gctgggacca caggcacaca ccactacacc cagctaattt ttgtattttt
                                                                   1560
attagagaca gggttteacc atgttagcca ggccagtctc gaactcctgg cctcaagtga
                                                                   1620
tetgeetgee teggeeteee aacatgetgg ggttacaage gtgageeact geacaggetg
                                                                   1680
cttgtttgtt ttctaacage catcctggag gggtgaggtg gtagctcact gtggtttga
                                                                   1740
tiggeactic cetegigaet tigtecatet tileaggige tiatigagea ticeigiati
                                                                   1800
ttccctggag aatgtcgtct tttcaacaac tttgcaccca cccccacctc cccgccaccc
                                                                   1860
cetetggitg tagagatggg gtettgatgt gittgeecag getgiteitt tgeecatitt
                                                                   1920
ttaattgggc tgctttctta ctgagttatg ggagttcttt ttatattctg gatatctatc 1980
ccttataagt atatgatttg caaatatttt ctcttaattt cccatatttc taagagacag 2040
tttcattaag taattaaaac acatacctaa attctgccg
                                                                   2079
```

<211> 1913

<212> DNA

<213> Homo sapiens

catttgcaga	tgctcctggc	aaagcatgtt	gttaagcact	atggtcagca	gatgaaattg	60
tctatgaaac	atcaactccc	caaaatgaag	acattccatg	aacctaccac	aattttgggt	120
aatagtttac	ctaaatgcac	tgaaattaag	ccagaagtta	acacattgac	tgcagagaat	180
aaattgtggg	atgatgcaaa	aaatggcttt	gcacggtgta	cagcigcgga	aatccaaaga	240
tttgcatttt	ctgctacagg	gctgttgtct	catgttgaag	agggtttgga	ticcgatgca	300
actgatagca	gctctgatga	cgatitggat	gaatatacce	ttagaaaaaaa	tgtggcagtg	360
taagtgcaaa	attattatta	gactattic	tgttccatat	atagcagcaa	ttatcttagt	420
ttccaggtat	gttgacaaga	aatagatttt	ctaaaatctt	aatgctataa	tettttttt	480
tttttttaat	ttttatttt	gagacagagt	ctcgctctgt	cgcccaggct	ggagtgtagt	540

```
600
ggtgcaatcc tggctcactg caacctccgc ctcccgggtt caaacaattt tcctgcttta
getteetgag tagetgggat taeaggtgtg tgeeaceaea eeeagetaat tittgtatit
                                                                  660
ttcgtagagg caaggtttca ccatgttggt caggctggtc tcgaactcct gaccttgtga
                                                                  720
tecacegee teggeeteec aaagtgetgg gattagagge gtgagecace acatecagee
                                                                  780
                                                                  840
accataatct tttatgttat aaaacttttg ttgaattttt ttaatgtttt gtttgttaaa
900
ggaaaaaccc atacaggaat aatgaaatta ttgagetata aataagcata ttttctattc
                                                                  960
ttgaataggc tgtggacaag gcctaatctt tgtttaagtg atctagttaa tatgtgtatc
                                                                 1020
                                                                 1080
taactaaaaa actttagtet geacataggg ageceteatt gtetttggga gtgtateagt
tgagagtaca tgtaagttga cttactactt tttttcctta actctctact cgtactcata
                                                                 1140
gettteagaa etgaeettta acaatteagt tagtttttge tagettagta taactaaaae
                                                                 1200
aaaactataa tgtcagctgt aagatatcta ttgaatgctt attatgtgct agacactaag
                                                                 1260
atteagtigi gageaacata ticacaacet eigeetittig gggeatgiae tigagagaga
                                                                 1320
ggtatetega tattgaataa taaaaageag agaaaaatag titeagitai cacaeegiga
                                                                 1380
taacactaca gaccaactet gteeaataga aacttetgag atgttggaaa tettttatgt
                                                                 1440
ctatgccatc taataggcac tagacttatg tggatattaa acacttaaga tttggccagt
                                                                 1500
gatactaagg aaatgagatt ttaattttat ttaattgact aaattttagt tgaaatggtc
                                                                 1560
agataaagca taatttttaa tttagttttc aggggatcta ttactgtccc caaattgatg
                                                                 1620
tgaattattg tttgtatata tagcattttg ggggaaagaa gtctgtcaca catggataca
                                                                 1680
tacaggggca caacactcac tggggctttt taaagggtgc agggtgggag gagggagag
                                                                 1740
atcaggaaaa ataactaatg ggcactaggc ttaaaacctg ggtgatgaaa taatctgtat
                                                                 1800
aacaaacctg catgacacag atttatctat gtaacaaacc tgcacttgta cccctgaact
                                                                 1860
taaaagttaa aaataaactt tiicaaaitc tcaaaaataa aigagaaita cag
                                                                 1913
```

<211> 2751

<212> DNA

<213> Homo sapiens

```
acteteaage gegeeggaa aggagggage agetteeggg acetggegeg gettitgtgt 60 tgggeagege gaatgtggeg ageteggtge gteteegetg eteetteee ttateeetgg 120 gaggteeaag tggteeegg geagettetg ttgetetggg acetgeaggt eeeggaaggt 180 eettagggag gaceeeagae aceggagaet gggaaatggg actattggea tteagggatg 240 tggetetaga atteteteea gaggagtggg aatgeetgga eeeageteag eggagttigt 300
```

```
360
atagggatgt gatgttagag aactacagaa acctgatctc ccttggtctt gctatgtcta
agccagaact gatcatctgt ctggaggcaa ggaaagagcc ctggaacgtg aacacagaga
                                                                   420
agacagccaa acactcagta gcgacgaggt ttcgccatgt tggccaggct ggtctcaaac
                                                                   480
teettacete aggtgateca cetgeettgg ceteceaaag tgetgggatt acaggcacgg
                                                                   540
gccaccactg ccagcctatt tgtgtattct gaattatatt taaccattca tttggtgagt
                                                                   600
tttgtcttct tatcttactg aagacatttt gccagagcag ggcctgcaag tttcattcca
                                                                   660
aaaagtgata ctgagaagat atgaaagatg ttgtcttgag aaattacgct taaggaatga
                                                                   720
ctgggaaatt gtggattatc cagactcagg tagttcttta taacaatgtg agaatgaact
                                                                   780
aatacagaaa agtggtacca gagagttggg acattgctat aaagatacct gaaaatgtgg
                                                                   840
                                                                   900
aagtgacttt ggaactgggt aacaggcaga agttggaaga gtttggaggg ctcagaagaa
gacaggaaga taaggaaaag tttggaactt cctagagact tgttgaatgg ttgtaaccaa
                                                                   960
aatgctgatg gtgatatgga caatgaagtc caggctgagg agttctcaga tggagatgag
                                                                  1020
gacettattg ggagetacag taaaggteae tettgetatg etttageaaa gagaetagtg
                                                                  1080
gcattgtgcc cctgctatag ggatctgttg aactitgaac tigagagaga tgatttaggg
                                                                  1140
tatetggcag aaaatattte taagtagcaa agcatteaag atatggcetg geteetteta
                                                                  1200
acagigtatg cicatattic tgaggaaaga gattatciga aaciggaaci tacgittaaa
                                                                  1260
1320
aaccattttc tggggaggaa ttcaacctag ctgcaaaaat ttgtgtaagt aaagaggagc
                                                                  1380
cgtatgttaa cagccaagac aatgggaaaa atgcccccaa gacatttcag agactttcgt
                                                                  1440
ggcaacccct ctcatcacag gcctggaggc ctaggaggga aaaacagttt tgtgggtcag
                                                                  1500
                                                                  1560
gettagggee etgetattet gtgeageett gggaeeetgt teeetgtget ttagetgete
cagetecage catggetaaa aggaetecag atatgtttea ggttgetget ecagagggta
                                                                  1620
taagacacaa geettggagg ettecagatg gtgttaagee tgeaggtget cagagggeaa
                                                                  1680
gagttgagge tigggageet ceattettte agattteiga ggatgtaigg aaacaacigg
                                                                  1740
atatccaggc agaaatttgc ttcaggggcg gagcccttgt ggagaacctc tactagggta
                                                                  1800
ctgtggaggg gaaatatggg gttgaagtcc ccacaaagag tctccactgg ggcactgcca
                                                                  1860
agtggagetg tgagaagagg gccactgtcc tccacacccc agaatggtag ctccatcaac
                                                                  1920
agtttgcact gtgtgcttgg aaaagccaca ggcactcaac accagcctgt gagagcggcc
                                                                  1980
atggggcact aagccetgca gagccgccag aagcagagct gtccaagacc ttgggagcct
                                                                  2040
accccttgca tcagtgtggc ctggatgtta gacatggaat caaaggatat tattttggag
                                                                  2100
ctctaagatt taatgactgc cctgctgggt ttcggacttg catggggcct gtaacccctt
                                                                  2160
tgttttggcc aatgtctccc ttttggaaca ggaacattta cccaatgcct gtacccttat
                                                                  2220
                                                                  2280
tgtatectag atgtaactaa ettgettttg attttacagg etcataggea gaagggaetg
ccttatctca gatgaaactt tggacttgga cttttgggtt aatgctgaaa tgagttaaga
                                                                  2340
                                                                  2400
ctttgggaga ctgtttggaa agcataattg tgttttgaaa tgtgaggaca tgatatttgg
gatgggccag gagtggaatg atatggtttg gctctgtgtc cccacccaaa tttcatgtca
                                                                  2460
```

aattgtaatc	ttcaatgttg	${\tt gaggagggtc}$	ctggtgggaa	ggtaattgga	tcatgggggc	2520
agacttctcc	tttgctgttc	tcatgatgag	tgagttctca	tgatacttga	tigittaaaa	2580
gtgtatagca	tttccccctt	tgctctctct	ctcctgccag	ccatgtgaag	atgtgcttgc	2640
ttcccctttg	ccttctgcca	tgattctaag	tttcctgagg	cctcccaga	agcagaagca	2700
tgtaaagccc	acagaaccgt	gagttgatta	aatctctttt	ctttataaat	t	2751

<211> 2816

<212> DNA

<213> Homo sapiens

```
60
atcitggcgg cggagcgatg agcgggtcta acccgaagge tgcggccgcg gcgtcggcgg
                                                                     120
ctgggcccgg ggggctggtg gctggcaagg aggagaagaa gaaggcgggc ggcggcgtcc
                                                                     180
tgaaccgcct gaaggcgcgg cggcaggcgc cccaccacgc ggccgacgac ggcgtcgggg
cagcggtcac ggagcaggag ctgctggcgc tggacaccat ccggcccgag cacgtcctgc
                                                                     240
                                                                     300
gcctcagctg ggtcaccgag aattatttat gtaaacccga agacaacatc tacagtattg
atttcacccg cttcaaaatt cgagatttgg agacagggac agtacttttt gagattgcca
                                                                     360
                                                                     420
aaccttgcgt ttcagaccag gaggaggatg aggaggagg aggtggagac gtggacatca
gcgcaggacg ttttgtccgc tatcagttca caccggcatt tctccgcctc cggacagtcg
                                                                     480
                                                                     540
gggctacggt ggagttcaca gtgggagaca aacctgtttc aaacttccgg atgatcgaac
ggcactattt ccgggaacac ttgctgaaaa actttgactt tgaltitggc tictgcalcc
                                                                     600
ccagcagtag gaacacttgt gaacatatct atgagtticc ccagciticg gaggatgtca
                                                                     660
                                                                     720
ttegtetaat gattgaaaat eettaegaga eeegetetga eagettetae tilgttgaea
                                                                     780
acaagctgat aatgcacaac aaggctgatt atgcctataa tggaggccag taagtgctgc
                                                                     840
aagagtaggt aggggaggtg ctttgccgcg gccacaagat cctggcacac ggagatgatc
gaagetgeag titgteaaca cacatetgga acetggeece aggaageeaa ggetggggtg
                                                                     900
                                                                     960
geagtiteet gegegeeaaa ggagetgeea aacagtgetg tgittiette eecagtatii
tttetteeet tttttteetg eeeegtaggt igeagaggta etaiagtaaa giaaaaggtt
                                                                    1020
                                                                    1080
aggataaggg teetggaate cagataaaaa agtttatttt eegtagttet ggetgeetgt
tggttgtctt gacgaccagg catagctgtg cctggtgaga aggctctggc caggcccatc
                                                                    1140
                                                                    1200
agcaggtcag cagcictiaa ggitcciggg igcigiggga agcigaaagg taggccicti
                                                                    1260
ccaggtaget ectectetea ceteeggeat tgeeateage geagtetgee eteggtetgt
gtgaagtett aaaccaactg gaagacactt gaaagggtgg ggagggaggg aggtgecaag
                                                                    1320
                                                                    1380
agtggaggca ccaaggaatg ggtgatgctg ccaagctgaa gggtctgctt tgtggagagg
```

```
ctgctgctct gtctgacttc cagggtctca gccagccctc ctgggaatag accaagtttt
                                                                    1440
cagcetggca gtgccttctg ttcccatttt ggaggacaga caagcttgct ccacatctcc
                                                                    1500
tggctcctcc cttctgagtc tcatgaaata gaatgagtca gctctgctca tggaacagta
                                                                    1560
                                                                    1620
gtatetettg aggecagage aggtettgta ttttgttttt ttatttecag aettettteg
                                                                    1680
gggaggtttt ataaaatgac agtggtgttc ccagcatatg tgatatgtgg ttagacttct
gatagtatca gcttccaggg gctaatctgg cttatgttgg gaggatatgc ttacgaatca
                                                                    1740
geageagett tetaaaggag agatttgaet tttetetgea etgeaeagee tggaggattg
                                                                    1800
                                                                    1860
gettttgatg gggatttgee teegaagete tttgtacatt tettgtttag gagggtttte
ctatctacct ttctactgaa gtagtttctg gaactttcct ggtggatcag agttacgtaa
                                                                    1920
                                                                    1980
tgcagtctga gccttcagac tgctagttag aattgtttta ggtgttcaga aagggcaaaa
taggetgatg tggcctgtca gagtgatgtg ttetcaaaaa agtteacttg cacatetgtg
                                                                    2040
ggctgctttt gtcctcagac ccttagtgga cagactccac aaaccctctg atgagacgat
                                                                    2100
tgatgtggcc agggtccagt tagcatcagt agaaggatgt cactaggaaa ggcccaggta
                                                                    2160
                                                                    2220
tctggtaagt gactgtgagg tgtcacagta cctgtgacag gagagtgtcc tgatgtgctt
gggagaaagg ccgtatgggg gccagggatg gaagagacag tgtgtggcca cagaaattcc
                                                                    2280
tgtccatcca ccaccagtgc tgctccctgt gtgggctcta gggcgagtgg ccccgaacct
                                                                    2340
tggcccagtg ctttgtccca ggccagagtc ttggcaatgc cacatgctgg cagctttctc
                                                                    2400
                                                                    2460
actgagaagg teetagetta eccetgtgtg etggeettgg atteageece gagagagggg
                                                                    2520
agagaccatt ceteetgtgg agtgggttee ttateaccag accggeeact etcagaactg
                                                                    2580
gegteeactg taaateeagg tgeettaegt gtggetetgt eeettatget geaggggaaa
                                                                    2640
getgeatige catigiteee acciecteae iggeagaaag aigceaggge igitageaci
                                                                    2700
gtetecteae ettetgttte teattgtgge teeteaaatg ggattigeat gtteetgtea
                                                                    2760
agegtaacaa caateeette tetetttgac agaggeeeag gtgggacagt ttetattatt
tgtataaaat gttattttgc cacatgagac agtaataaaa gaaagatttt cacagt
                                                                    2816
```

<211> 1766

<212> DNA

<213> Homo sapiens

acttgaggtc	ggtgtgggga	acttgctttt	aattctcatt	tagagaagac	agtactgaaa	60
tggagaaaag	tcacagggaa	agtactttta	cagattgtag	attagtaaag	aacccaaaga	120
gagcctttca	ttgagagcag	aaaggcgaat	ggaattcgct	gttttctgtc	taaggaggag	180
gaggatgggc	aggcaggtca	gctgcccagt	ggggcttggt	gtgatagtgg	gagtcaccct	240

```
teattigaac etetetgeet tgeeeagete eagticaget teagegtggt eagagaeact
                                                                     300
atctctatgg aaggtcactc ctggaagaat acatttactt agctgcttcc accatggaat
                                                                     360
cclagetigt geiggagigt eccetteate electeeigt geitigagaa tecatigiig
                                                                     420
ctggtatgcc ctgagcagtg cccttgaact tgcccaggta ccccttgaca tccacaccac
                                                                     480
aaatagteta geettacaaa ggtggacaag atgtetttte aacagtetgt aetgecaett
                                                                     540
ccatccatct gaagctiict giiccigagi cigicalgac attaatciit caaaaaicii
                                                                     600
tcacagagat tittagicic tactaaaaat taccaaatgc tictaaatat gaaggagagg
                                                                     660
                                                                     720
ltggggacac gcaccctatg tgataccaag ttttattgtc aagacagtgt catggtgcag
                                                                     780
aggtaggcat totgagcagg ggaacaaaat aagggcotag aaactcacco gtgcatatgt
tgacctttgc aaaatgacct ggtgacatgg caagtcagtg gggacaggaa ggaccactcc
                                                                     840
ctaagtaatc ccagaacaat ggctattcat gtgggaaaaa aagaaatttt actttctct
                                                                     900
accttacctg gtgataagtt ccaaalatgt taagggcttt aatacaaaaa gcaaaaattg
                                                                     960
tcagtgtttg gatgaaaaaa gccttagggc aggaaagaat ctcttgagac ataaagtagt
                                                                    1020
aatcalaaag gacaagalgg ttaagtcaat tctgttaaaa ctcaaggctt atattaagca
                                                                    1080
aacactigaa gigagaagat gatccacaac tigagaagac attiataata caaataactg
                                                                    1140
                                                                    1200
atgaaggatt cataatcaca aatatagaga attcctattt aaaaaaatag aaaaatagtg
aagactacac aagaggaaat agggctttta aataaataga tgttctgtag cattggtcag
                                                                    1260
                                                                    1320
ggaaatatga attaggacca caatgagatt ccattttata tccataagat ttgcaaaggt
tgggtctgac agtaccagtt gttagatctg tagggacttg tacaacattg tggatgtgta
                                                                    1380
aacaggcacc actgctttaa aaaacaattt tcccttacag acttgaacat ttgcagacgt
                                                                    1440
talgatelig eliceaacte ceaceigial giceageaaa etetigeaig iggeeaciag
                                                                    1500
gaggaatgtg taagaatgtt catagttaca tatttataat agttaataac tggaaaaagt
                                                                    1560
gaaatgtatg tctgtctaca ggaaaatagg tgaataatta gatatatata ttcattctac
                                                                    1620
                                                                    1680
gggalattat tcagtagtgg aaatgagtga actacagcta tacctcacaa taagaatgaa
tctcagaaaa tattaaggaa aaaagcaagt ttgaagagac cacatggggc gtactatttt
                                                                    1740
tattgagccc aaaaacaagc aaaacc
                                                                    1766
```

<211> 3359

<212> DNA

<213> Homo sapiens

<400> 2058

aaatctacct atagtccttg titctggagg tigttgccat ggigagatti gatticatgt 60 atgttctitt giggtctatt aacctagcca tcatcattga tittattatt titgagtcag 120

agtcgcactc	tgttgctcag	gctggagtgc	agtggtgtaa	tcttggctcg	ttggaacctc	180
cgcctcccag	gttcaggtga	ttcttgtgcc	ttagcctctg	gagtagctgg	gattacaggc	240
acgcaccacc	atgcctggct	acttttgtaa	ttttagtaga	gacggggttt	cgccgtgttg	300
gccaggctgg	tcttgaactc	tggcctcaag	tgatctacct	gtctcagcct	cccaaagtgc	360
taggattgta	ggagtgagcc	actgtgcctg	gcctggtttt	attattacta	tttttaatat	420
ttgttttttc	atatgataga	gacagtgtct	tgttatgttg	cccaggctgg	tcttcaactc	480
ctgggctcga	gatcctcctg	cctcaacctc	ccagagtgtt	ggtattatag	gcgggagcta	540
ccgtgcttgg	cccagtttta	ttattttaaa	atagtaagtt	agccattaca	cttaagatgt	600
gaaaattcca	aatatagtgt	taaaaaagta	catagaagac	tgatttttcc	ctttctgaaa	660
ctgtagagaa	gcagttttct	aggccatgaa	aaaacggcaa	gagccttatt	aaatatataa	720
tttgaagcat	ttttaaatat	agatttgatt	ggagatagaa	acttggccaa	gctgttacta	780
ctccatctta	taggcagaat	aataatgtga	tttctcaaaa	taaaaataga	aaagcaaaaa	840
ctgggtcttg	ctgctagaaa	accagcttcg	agattggctt	catgttttca	aaatcctgat	900
aaatttaata	ttgatgtccg	cgaagtattc	attigitgaa	taaattaatt	tgagcaaaaa	960
ttatatttta	gttatattta	catttttaaa	ataaaataga	aaaatccctt	attaccctgc	1020
ttctccaaat	agctctgtta	atttgtgcat	atttacttta	agttttttgt	agttgcagtc	1080
actaatatcc	agactgcttt	gaattctggt	ttggaaaaaag	ctcagtattg	taaacctttc	1140
ctcatgtttt	tgcagggcct	ctacttttgt	tgactgtaaa	tttttcaaca	gtcatgctga	1200
tgtcctaatg	acctgcttgt	ttttggtgga	tttacttagt	gggagcagga	gctgaggtta	1260
tgcgtgttta	gtcctccagc	cttgaaattc	ttacagcctt	tcagggactc	agtactgatg	1320
tgactgaatt	ggacttgaag	agtagatttc	ctttgtgtga	attaggtgga	actgtttatg	1380
catgictggg	ttgctaaagg	gaaaggaagt	gagttgagaa	gggaagggag	acatactttt	1440
gtccaaattt	atgccctaac	agtctgattt	tttttttga	atatagaaat	acttgttaaa	1500
tatcttccat	caacagataa	acagatggac	aaaaagattt	ctattttaaa	ggatcatggc	1560
tatatagaaa	atttgacatt	tggatgggat	ggaccatctt	ggaggctact	cacagccctt	1620
aagttgttat	gtctggaagc	tgagaaattt	acatgctgga	aaaaagtact	tcttggggag	1680
gtaatttcag	atacgaatga	gaagacaagt	ttggacatag	cccagaaaaat	atgctattat	1740
ttcatagaag	agactaatgc	tgtgcttcaa	aaggtgtctc	atatgaagga	tgaaaaagag	1800
gccctgataa	accaactaac	tttggtggaa	tccttgtgga	cggaagagct	aaagattctc	1860
agggcatctg	ccgagaccct	gcacagtttg	caaacagctt	ttacctgatt	tcaccgaagc	1920
gcatttggtc	acctcctctg	aaacaaaagt	taattttgaa	gagcatcatc	atgggctggg	1980
gtggtggctg	ccccaggac	atgcaggatt	tctgcagggg	gcagcacagg	ttctgggatt	2040
gtgaggctgt	gagtgaaggt	ggacaagctg	tctggatggc	aggtctaatg	ctcttccgaa	2100
taaagtgctg	aactgtgagg	agagaggcgg	actgtgaggc	agccaggagc	cagctgcgtc	2160
cgtgtgtggt	ctgtcaccac	ggggcctgct	tcttatctga	cacagcagct	atcagagtct	2220

agtggttgtg	cttttaagat	gctctgatac	cattgggtta	aggggcagat	tggcggtggg	2280
tgtggggcag	tgtgaggtag	tcctggatcc	ccgccagggt	ggcccagacg	ccagcccttc	2340
cctgtgtggc	tgcactgagg	tgggtgttga	agagccccct	aggggacaca	cagcttccag	2400
gaggagggaa	tgtcctctaa	gcatgctcct	ggcctctcaa	ggtggcgctt	gtctaattat	2460
tcacttggga	agaatgacta	gctcagccag	cggctctttc	tgctttgttc	tggcgacttt	2520
cctgggcagg	cctttccacc	tggggagctg	gctcatcctg	cacagctggg	ccgtggtggg	2580
cctgtctgct	tgattctggg	gttcagtgta	ggtcagctga	tggcgaacca	tggtggtggt	2640
ttggcttctg	ttcttattct	tgagttttga	taccacgcag	accttgggtg	gggagagctt	2700
cctgcacagc	tctcagcggc	ctgtggcctt	ggaactgcct	gcgtaagtaa	cggaggggct	2760
gctggtcctg	ttcaggcccg	tgctggggac	gccgcttaga	caatgttgcc	cagagtcctg	2820
tttaccctcc	cagggttcat	tcttcccaag	aactcaaatt	cctttctcat	tggagcctag	2880
tgaaaccaaa	tgaacgggac	ctgctggcct	caggaggcag	gcagagttta	aaataaaact	2940
ttctcatgat	ttcttgaaca	tctttccctg	tttgtatata	cactttgtgt	ttatttttca	3000
gtagctgcag	tatattttt	ttcaatattc	agtataatgc	agtgtatttc	atcatatgct	3060
gtatggagag	tgggcagact	tctgtggagg	gcccgatagt	aaccatttga	agctttctgg	3120
acctgtggtc	ttagtcccag	cgattctgca	gagcggccat	cggcagcatg	tcaaccattt	3180
gcatggctgg	gctccaggga	aactactgac	aacgacaggt	ggtgggccat	agtttcctga	3240
cccctgtgct	atgccagaat	ttcttttcc	tcttccctat	gagtggacct	aaatatgtta	3300
attccttttc	acctttcaaa	acggacagcc	ccttgaacat	taaaaacttt	gcagaccct	3359

<211> 1692

<212> DNA

<213≻ Homo sapiens

tcaagccaga t	gtctcacta	tgagacaact	gctcagccag	cccagaagta	aaacaatgtg	60
tctgaaatgt g	gatetecaag	agcgactgct	ctgcccatcc	ctactcgctg	gcacagctga	120
cggctccttg a	ngaatggatg	accctaaagg	agacttcatc	acactctacc	agatggcttc	180
ccagtcatcg g	gcctctcatt	acaageteca	agtgatcaag	gctttaaaat	ctagcgggct	240
cigcgagtca t	tgacatatg	gactcccgtt	catcctcaga	cctacaagct	gttggcagct	300
ggactgggat g	gagctggaga	caaatcagca	acatttccat	gctttgtgtc	acagcctgct	360
gaaaagggaa t	ggctgctgt	tagccaaggg	ggaaccaccg	ggcccaggac	acagccagag	420
aattcctgcc a	ngcaccttct	atgtgatcat	gccgtcacac	tccctcacac	tgctggtaaa	480
ggcggtggcc a	cgcgggaac	tgatgctgcc	cagcaccttc	ccctgctac	ctgaggaccc	540

acatgatgat	agccttaaga	atgtggagag	catgctggac	agcctggagc	tggagcccac	600
ctacaacccc	ttgcatgttc	aaagccacct	gtactcacac	ctgagcagca	tctatgccaa	660
gcctcagggg	cggctccacc	cacactggga	gagccgagct	ccgagaaaga	ctgggcagtt	720
gcagaccaac	cgagctcgag	ctactgtggc	cccctgcct	atgactcctg	tcccaggcag	780
agcctccaag	atgccagcag	ccagcaaatc	ttcctcagat	gccttcttcc	tgccttcaga	840
gtgggagaag	gatccctcaa	ggccctaagt	caccagcacc	agagcccagc	tgcccagctt	900
aaccatatcc	atgctcaggt	tcacataatg	gctatctgtg	gtcagacttg	ctctctatcc	960
gcctgagcct	ctgtgagtga	gggctgactg	ggaaacaaca	gccttcctgt	cctgtttcag	1020
tgctgtccca	ctcctcaagt	ctggaagcga	cacacccgag	cctgtccttt	ctccagcaag	1080
gactttcatt	ttctttagaa	tcatttgcta	ctgtttacac	aggtgaagat	taaacaccca	1140
gtaagcttct	accattgtta	ggagcattca	taactcagaa	tttcttcttg	tagctctgtg	1200
taagcaggtg	gatgaggtca	gatcaccttt	ggtaaactgg	acctcaggaa	caaggatgag	1260
gttttgaaag	ctcataaaaag	acaagtaaga	ttgaaatcca	agcctcattt	cagagcctgt	1320
gcccttccca	ctacaccacc	aggetteage	ctccaaagag	acaagtgctt	ggtacctaca	1380
tgcaaagtgt	gtgtgctggg	gggtgggagg	gctgcccaga	acaggggaga	ggatggtgta	1440
aaaaaagacc	tactcctttc	ctgttaccct	ctccccacat	gtaccaacct	tcctgttgct	1500
ccctccatcc	acagaataat	agctaccatt	tataaaatgt	ttactctggg	ctgggagcag	1560
tggctcacac	ctgtaatccc	aacactttga	gaggctgagg	tgggatgatc	acttgaggcc	1620
aggagttcga	gaccagcctg	agcaacactg	tgagaccccc	ccgccatctc	tacataaata	1680
ataaaaactt	tt					1692

<211> 2269

<212> DNA

<213> Homo sapiens

aggcgcgcgg	gaacatgggg	ctgtatgctg	cagctgcagg	cgtgttggcc	ggcgtggaga	60
gccgccaggg	ctctatcaag	gggttggtgt	actccagcaa	cttccagaac	gtgaagcagc	120
tgtacgcgct	ggtgtgcgaa	acgcagcgct	actccgccgt	gctggatgct	gtgatcgcca	180
gcgccggcct	cctccgtgcg	gagaagaagc	tgcggccgca	cctggccaag	gttcatcggg	240
gtgtgagccg	gaatgaggac	ctgttggaag	tgggatccag	gcctggtcca	gcctcccagc	300
tgcctcgatt	tgtgcgtgtg	aacactctca	agacctgctc	cgatgatgta	gttgattatt	360
tcaagagaca	aggittcicc	tatcagggtc	gggcttccag	cctcgatgac	ttacgagccc	420
tcaaggggaa	gcattttctc	ctggacccct	tgatgccgga	gctgctggtg	tttcccgccc	480

```
agacagatet gcatgaacac ccactgtacc gggccggaca cctcattctg caggacaggg
                                                                      540
ceagetgtet eccagecatg etgetggace eccegecagg eteccatgte ategatgeet
                                                                      600
gtgccgcccc aggcaataag accagtcact tggctgctct tctgaagaac caagggaaga
                                                                      660
tctttgcctt tgacctggat gccaagcggc tggcatccat ggccacgctg ctggcccggg
                                                                      720
                                                                      780
ctggcgtctc ttgctgtgaa ctggctgagg aggacttcct ggcggtctcc ccctcggatc
caegetacea tgaggteeac taeateetge tggateette etgeagtgge tegggtatge
                                                                      840
                                                                      900
cgagcagaca gctggaggag cccggggcag gcacacctag cccggtgcgt ctgcatgccc
tggcagggtt ccagcagcga gccctgtgcc acgcactcac tttcccttcc ctgcagcggc
                                                                      960
                                                                     1020
tegictacte caegigetee etetgecagg aggagaatga agaegiggig egagatgege
tgcagcagaa cccgggcgcc ttcaggctag ctcccgccct gcctgcctgg ccccaccgag
                                                                     1080
geetgageae gtteeegggt geegageaet geeteeggge eteeeetgag accaeaetea
                                                                     1140
gcagtggctt cttcgttgct gtaattgaac gggccgaggt gccaaggtga gtgagtgggg
                                                                     1200
gcgtgcttgg gaggcgcagg atggcaccgg cacatctaac atctacactt etctagetca
                                                                     1260
gcctcacagg ccaaagcatc agcaccagaa cgcacaccca gcccagcccc aaagagaaag
                                                                     1320
aagagacage aaagageege ageeggtget tgeacaeege ettgeacata geagaggete
                                                                     1380
                                                                     1440
egggetgaet cetteetggt gggaaaggaa gatgeetgte eteteegtgg aggaeeetgg
gccctcaccg caggaagcag tttgggtttt gaaaggttat tgggtccctt ccttgggctg
                                                                     1500
tgttcttgct ggtgagcaaa gtgttgcctg caaaaataaa atgcagaacg tactctacga
                                                                     1560
tagatcacag ttttttattc ttaatgtcac aagcaggaga aaaatctcac attcatacta
                                                                     1620
aaagttecaa etagacteaa caggaatgaa gtetetattt gtaatggaaa gteecageet
                                                                     1680
ecegelgeeg tecagtgegt gtaetgtaca catecacact cacacteact cagggttece
                                                                     1740
ggaccggctg teetgeetge ggaactgagg taaactaget caggtgetga cactaggagg
                                                                     1800
gtetacetta cataaggtae aggtagaage ttgattgeta ggeecaggee cacceagace
                                                                     1860
                                                                     1920
ctccaatect aacgggtatt taggettgag gttcactece tectcagetg cacaegeage
caggtattaa cgaggatcag agctgttctg aggggtggga aggagcagcc ccaccaccac
                                                                     1980
teacteacce teagteacat eggggagggg geaceagtta catttacate acattattta
                                                                    2040
taaaataaga attacattte atataacatg gecagaagga getetagtee eecaggaaag
                                                                    2100
\verb|ctgccgggga|| cagcatttga|| gcctcttctt|| tgcacaggca|| tgacttaact|| atacagctaa||
                                                                    2160
tteetagtta atageattta taettaaeca eeteaatgaa eeaagettga aggaatttaa
                                                                    2220
aaggcaattt agcttaaata caaaaataaa tttttgttaa aaaacgttt
                                                                     2269
```

<211> 2395

<212> DNA

<213> Homo sapiens

aagtcaggac	gggagtccgg	cgggttacag	cggaggccta	ggtggcagac	agggggcccg	60
ggccgctgcg	tgttgtccac	ccaagatgga	gttcctcctg	gggaacccgt	tcagcacacc	120
agtggggcag	tgcctcgaaa	aggcaacaga	tggctccctg	caaagtgagg	attggacgtt	180
gaatatggag	atctgtgaca	tcatcaatga	gacggaggaa	gggccaaagg	atgccattcg	240
agccctgaag	aagcggctca	acgggaaccg	aaactacaga	gaggtgatgc	tggcattaac	300
agtgctggag	acatgtgtga	agaactgtgg	ccaccgcttc	cacatccttg	tggccaaccg	360
agatttcatc	gacagtgttc	tggtcaaaat	tatatctccc	aagaacaacc	ctcccaccat	420
tgtacaggac	aaagtgcttg	ctctgatcca	ggcatgggct	gatgcctttc	gaagcagtcc	480
tgatctcacc	ggcgttgtgc	acatatatga	ggagctgaag	aggaaggggg	ttgaatttcc	540
catggcagac	ttggacgctc	tgtctcccat	acacacacca	cagcggagtg	tccctgaagt	600
ggatccagct	gcgaccatgc	ccaggtccca	atcacagcag	aggacaagtg	ctggttccta	660
ttcctcgccg	cctcctgctc	cctactccgc	accgcaggcc	ccagctctga	gtgtgactgg	720
ccccatcaca	gccaattcag	aacagattgc	caggctgcgg	agtgaactgg	acgtcgttcg	780
aggaaacaca	aaagtcatgt	ctgagatgtt	aacagaaatg	gtccctggac	aggaggattc	840
atctgatctg	gagttgctgc	aggagctcaa	caggacctgt	cgggccatgc	agcagcgcat	900
cgtggagctc	atctcccgcg	tgtccaatga	ggaggtcacc	gaggagctgc	tgcatgtgaa	960
cgatgacctc	aacaacgtct	tccttcgata	cgagaggttc	gaacgataca	ggtctggccg	1020
atccgttcaa	aatgccagta	atggagtact	gaatgaagta	accgaagaca	acttaataga	1080
cctggggcca	gggtctccag	ccgtggtgag	cccaatggtg	gggaacacag	cgccccatc	1140
ttccctctcc	tcccagcttg	caggcttaga	cttggggaca	gagagcgtca	gtggcaccct	1200
cagttcactc	cagcaatgta	atccccgtga	cggctttgac	atgtttgccc	agacgagagg	1260
aaactccttg	gctgagcagc	gcaagacggt	aacctatgag	gatcctcagg	ctgtcggagg	1320
acttgcttct	gcactagaca	atcgaaaaca	gagttcagaa	gggatccccg	ttgcgcagcc	1380
atctgtcatg	gacgacattg	aggtgtggct	caggaccgac	ctgaagggtg	atgatctgga	1440
ggagggtgtc	acaagtgaag	agtttgataa	attccttgaa	gaaagagcca	aagctgctga	1500
aatggttccc	gacctcccct	cgcccccat	ggaggctcct	gccccagcct	caaacccttc	1560
tggccggaag	aagccagagc	ggtcagagga	tgccctcttc	gccctgtgag	cagctctgtg	1620
gtttgcctcc	ccagatggcg	ggtccccgct	cgcaccccgt	ggacaccggg	cactggccac	1680
tcctacatcc	ccagctccac	acggcctgca	cacctgtgtt	tccatggaaa	tgccaccgtg	1740
tctgctccca	ggcctcccac	tagtcaggac	cagcttcagc	cacttcttt	ctctgagtgg	1800
tgggacaact	gcagccagag	actctctccc	ctcccaccat	gggcccctct	gcccatgttt	1860
cctcccagga	agagcgggca	gagtggccca	gccccaggca	gtgcttcctg	agcagaccac	1920
ccggactgtc	tttcctccac	ccgcccatgg	agaaagagca	cgcccggccc	cgccctgtgc	1980
teacctetge	ctggctcagc	gaccttctca	ggcattctgc	cctcctgggc	ccctctctcc	2040

ctgaaggggc	tttgtggcat	ctctggaaga	gcagggtgtg	${\tt ctgcactcat}$	gggcctggtc	2100.
tcactccttg	gacttgtcac	cttgtgacat	ttggcttatc	agcatttgag	aaggctctgc	2160
tgggtctcca	tggtgggggt	ctctcacctt	cttgaccctc	tctccatcat	tcagctgcca	2220
gcccaggctt	cacacccaag	ctggctcagc	agccgagcct	ggcaccgagg	gtccctgcag	2280
gctccctggg	cagggagagg	gccaaggaca	attgggaggg	cagcaggcag	cccgcagatg	2340
gtggccatgt	ggcacgctgc	tgagacgaca	ctaccaataa	accaaactgc	cacgc	2395

<211> 2284

<212> DNA

<213> Homo sapiens

acggggccgc	ctggagaggt	gctgggagct	gggtggagct	tagaggaatt	aaactttggc	60
cctgcgcctc	gtccagccta	ggttccaccc	ttttctggga	acaatgaatc	tcgctgtgtt	120
gtccaggctg	gagtgcagtg	gcaccatctc	ggctcactgc	aacctctggc	tcccaggttc	180
aagcgattct	cctgcctcag	cccctgagt	agctgggatt	acaggcacgc	gccaccactc	240
ccaggctccg	gtagattgca	aatgacctgc	tttctttctg	ttcccgggcg	tttggacccc	300
tgtcttggac	cgctgtcgga	tagtaaatcc	caagtaaggt	acctgccgtc	ggcagatttg	360
agctttcttc	ttggacacct	aatacccaga	gtcctccagg	ctccggtaga	ttgcaaatga	420
cctgctttct	ttctgttccc	gggcggcatc	ggacccgtcg	gagagtaaat	cccaagtaag	480
gtacctgccg	ttggcagatt	tgagctttct	tcttggacac	ctaataccca	cagtcctcca	540
ggctccggta	gattgcaaat	gacctgcttt	ctttctgttc	ccgggtggca	tcgacccgtc	600
ggagagtaaa	tcccaagtaa	ggtacctgcc	gttggcagat	ttgagctttc	ttcttggaca	660
cctaataccc	acagtcctcc	aggtgagtcc	taaggatett	aggatacgcg	atgggggtcc	720
taaggcaggg	ggggaagagg	ggatggctgt	cacccaaccc	aaaatgggcg	gcctttatgt	780
tcaggttttg	cccaagagtc	agcttatttg	cttcttgtac	tatcagggca	gttgatgcca	840
cggccctcaa	acatgagggg	ccatccttta	gaaaccctct	ctagttgttt	agacaactag	900
gccaccggcc	tcagccaggg	ccccagagtt	teggttaaaa	gtccagctgc	catcttttct	960
ctatctgacg	cattcaatgg	aaaaggcttt	gtcagatcgg	gtagccccag	ggctggggct	1020
gecagaagtt	tttcctttaa	ctcctgaaag	actttttgtt	cttgggatcc	ccattccaaa	1080
ggitccgitc	cccgccccct	ttgtgacctc	atacaaaggc	ttggctaata	ctgcaaagtt	1140
tgggatccag	tctacaaaac	cacacagete	ccaagaattc	ccttacctgc	cttctgccct	1200
taggctccgg	tagattgtaa	ataacctgct	ttctttctgt	tcccgggctg	cgttcggacc	1260
ccigtcggat	agtaaatccc	aagcaaggta	cctgccgtca	gcagatttga	gctttcttct	1320

tggacaccta	atacccacag	tcctccaggc	tccggtagat	tgcaaatgac	ctgctttctt	1380
tctgttcccg	ggctgcgttc	ggacccctgt	gggatagtaa	ctcccaagta	aggtacctgc	1440
cgtcggcaga	ttggagcttt	cttcttggag	acctaatacc	cacagtcctc	cagaaaaaaca	1500
aacaaagaca	tggatttact	gtgcatatta	gcagatccat	actggaaaat	gcatggaggt	1560
ttcatataca	ccacttacag	ttttcagctc	ctcagtagtg	acaaagccat	acccatcatt	1620
gtcgattcga	tcaacaatct	tccctagcct	ctcctcgctc	tcgtccgggg	tgagctcgtc	1680
gaagttcttg	gagtccttct	tgcccaggaa	ggcctcgtgg	tcgtactgga	agctctggtt	1740
gtcctcaggg	ggccgctcgc	ccagctccga	gtcgggccgc	accacgcgct	ctttgcgcac	1800
cgtgggcttg	gcccgcagaa	cccgcggcgc	cagcaccagc	gccagcagca	gccccagggc	1860
taaccccggc	ggccaccgcg	cgccatcgtc	ccgaggagag	ggcggccggg	agggagacgc	1920
tgagcgagcg	acaacagcgg	cagctcggga	atgggggctc	ggagcgcggc	ggccaagttt	1980
tatgttatgt	atatttaca	agtaaaaaaa	ttttttcacc	tcagcctgaa	ctgaacacta	2040
gctgacagac	gttttgattt	ctitgaccat	cacggaatcg	tggccaagcg	cggtggctca	2100
catctgtaat	cccaacactt	tgggaggtca	agatgggcgg	attgcttggg	tccaggtgtt	2160
tgagatcggc	ctgggcaaca	tgacaaaacc	ctgtttctag	taaaaataca	aaaattaacc	2220
aggetcaage	catgaccatg	caccattgca	ctccagccta	ggcgacagag	caggaccctg	2280
tctc						2284

<211> 3914

<212> DNA

<213> Homo sapiens

```
gaagagaaag aaaggactgg ctgggttgta ggcagcaggg ccgagcagct gagggctaag
                                                               60
120
tttagttcaa catgggctgt atccgaatcc ttctgaaatt tgctgggatt ccatgaggga
                                                               180
gleagglaca ecaaacegel cacelliget gacigcalla gigalgagii geegelagga
                                                               240
tgggaagagg catatgaccc acaggttgga gattacttca tagaccacaa caccaaaacc
                                                               300
acteagaltg aggateeteg agtacaatgg eggegggage aggaacatat getgaaggat
                                                               360
tacciggigg iggcccagga ggclcigagi gcacaaaagg agaictacca ggigaagcag
                                                               420
cagegeeigg agetigeaca geaggagtae cageaacige aigeegieig ggageataag
                                                              480
                                                               540
cigggciece aggleageti ggieletggi teateateea geleeaagia igaecetgag
atcctgaaag ctgaaattgc cactgcaaaa tcccgggtca acaagctgaa gagagagatg
                                                              600
giteacciec ageaegagei geagiteaaa gagegiggei iteagaeeei gaagaaaate
                                                              660
```

```
gataagaaaa tgictgatgc tcagggcagc tacaaactgg atgaagctca ggctgtcttg
                                                                     720
agagaaacaa aagccatcaa aaaggctatt acctgtgggg aaaaggaaaa gcaagatctc
                                                                     780
attaagagee ttgccatgtt gaaggaegge ttccgcactg acagggggte tcactcagae
                                                                     840
ctgtggtcca gcagcagctc tctggagagt tcgagtttcc cgctaccgaa acagtacctg
                                                                     900
gatgtgagct cccagacaga catctcagga agcttcggca tcaacagcaa caatcagttg
                                                                     960
gcagagaagg tcagattgcg ccttcgatat gaagaggcta agagaaggat cgccaacctg
                                                                    1020
aagatccagc tggccaaget tgacagtgag gcctggcctg gggtgctgga ctcagagagg
                                                                    1080
gaccggctga tccttatcaa cgagaaggag gagctgctga aggagatgcg cttcatcagc
                                                                    1140
ccccgcaagt ggacccaggg ggaggtggag cagctggaga tggcccggaa gcggctggaa
                                                                    1200
aaggaccigc aggcagcccg ggacacccag agcaaggcgc igacggagag gitaaagita
                                                                    1260
aacagtaaga ggaaccagct tgtgagagaa ctggaggaag ccacccggca ggtggcaact
                                                                    1320
etgeacteec agetgaaaag teteteaage ageatgeagt eeetgteete aggeageage
                                                                    1380
cccggatece teaegteeag eeggggetee etggttgeat eeageetgga etecteeact
                                                                    1440
leageeaget teactgacet etactatgae ecettigage agetggaete agagetgeag
                                                                    1500
agcaaggtgg agttcctgct cctggagggg gccaccggct tccggccctc aggctgcatc
                                                                    1560
accaccatee acgaggatga ggtggccaag acceagaagg cagagggagg tggccgcctg
                                                                    1620
caggetetge gtteeetgte tggeacceea aagteeatga ceteeetate cecaegttee
                                                                    1680
totoloct coccoccc accotytic colotaty otgaccoct cotygotyt
                                                                    1740
gatgccttcc tcaactcctt ggagtttgaa gacccggagc tgagtgccac tctttgtgaa
                                                                    1800
ctgagcctig giaacagcgc ccaggaaaga taccggctgg aggaaccagg aacggagggc
                                                                    1860
aagcagcigg gccaagcigi gagtacggcc caggggigig gccigaaagi ggccigigic
                                                                    1920
teageegeeg tateggaega gleagtgget ggagaeagtg gtgtgtaega ggetteegtg
                                                                    1980
cagagactgg gtgctlcaga agctgctgca tttgacagtg acgaatcgga agcagtgggt
                                                                   2040
                                                                   2100
gegaceegaa tteagattge eetgaagtat gatgagaaga ataageaatt tgeaatatta
atcatccage tgagtaacet ttetgetetg ttgcageaac aagaccagaa agtgaatate
                                                                   2160
egegiggeig tecticetig etetgaaage acaacetgee tgiteeggae eeggeeteig
                                                                   2220
gacgecteag acactetagt giteaatgag gigiteiggg tatecatgie etatecagee
                                                                   2280
                                                                   2340
etteaceaga agacettaag agtegatgte tgtaceaceg acaggageca tetggaagag
tgcctgggag gcgcccagat cagcctggcg gaggtctgcc ggtctgggga gaggtcgact
                                                                   2400
egetggtaca acetteteag etacaaatac ttgaagaaac agagcaggat gtitteaceg
                                                                   2460
agaaageete acctgatatg gatgggtace cagcattaaa ggtggacaaa gagaccaaca
                                                                   2520
cggagaccèc ggccccatcc cccacagtgg tgcgacctaa ggaccggaga gtgggcaccc
                                                                   2580
cglcccaggg gccattlett cgagggagca ccatcatecg ctctaagace ttctcccag
                                                                   2640
                                                                   2700
gaccccagag ccagtacgtg tgccggctga atcggagtga tagtgacagc tccactctgt
ccaaaaagcc accttttgtt egaaactccc tggagcgacg cagcgtccgg atgaagcggc
                                                                   2760
cliccleggt caagtegetg egeteegage gietgateeg tacetegetg gaeetggagt
                                                                   2820
```

tagaccigca	ggcgacaaga	acctggcaca	gccaattgac	ccaggagatc	tcggtgctga	2880
aggagctcaa	ggagcagctg	gaacaagcca	agagccacgg	ggagaaggag	ctgccacagt	2940
ggttgcgtga	ggacgagcgt	ttccgcctgc	tgctgaggat	gctggagaag	cggatggacc	3000
gagcggagca	caagggtgag	cttcagacag	acaagatgat	gagggcagct	gccaaggatg	3060
tgcacaggct	ccgaggccag	agctgtaagg	aacccccaga	agttcagtct	ttcagggaga	3120
agatggcatt	tttcacccgg	cctcggatga	atatcccagc	tctctctgca	tgacgtctaa	3180
tegecagaaa	agtatttcct	tigitccact	gaccaggctg	tgaacattga	ctgtggctaa	3240
agttatttat	gtggtgttat	atgaaggtac	tgagtcacaa	gtcctctagt	gctcttgttg	3300
gtttgaagat	gaaccgactt	tttagtttgg	gtcctactgt	tgttattaaa	aacagaacaa	3360
aaacaaaaca	cacacacaca	caaaaacaga	aacaaaaaaa	accagcatta	aaataataag	3420
attgtatagt	ttgtatattt	aggagtgtat	ttttgggaaa	gaaaatttaa	atgaactaaa	3480
gcagtattga	gttgctgctc	ttcttaaaat	cgtttagatt	ttttttggtt	tgtacagete	3540
caccititag	aggtcttact	gcaataagaa	glaatgcctg	ggggacggta	atcctaatag	3600
gacgtcccgc	acttgtcaca	gtacagctaa	tttttcctag	ttaacatatt	ttgtacaata	3660
ttaaaaaaaat	gcacagaaac	cattgggggg	gattcagagg	tgcatccacg	gatcttcttg	3720
agctgtgacg	tgtttttatg	tggctgccca	acgtggagcg	ggcagtgtga	taggctgggt	3780
gggctaagca	gcctagtcta	tgtgggtgac	aggccacgct	ggtctcagat	gcccagtgaa	3840
gccactaaca	tgagtgaggg	gagggctgtg	gggaactcca	ttcagtttta	tctccatcaa	3900
taaagtggcc	tttc					3914

<211> 5245

<212> DNA

<213≻ Homo sapiens

tccctgttgt	tctaaattcg	gcattactag	tgcatgcgtg	catccgggga	aaaggaacaa	60
ggtgggagaa	gagagagaaa	gegaatacce	gaggeegeea	gcatcagtgg	gtgcccgcgc	120
tetectecte	gctctcgtcc	tetgeeetee	gecetggete	cctgcccgca	ttccctggga	180
gcgcagcctt	gccttagcct	gggagacagc	tgtccacagt	gacaggcggc	cattgttctc	240
ggccgagcca	gcaggcttcc	ggccggtggc	agctgctgct	cctccgctct	geggeeceae	300
caagggggcg	ccgccaccgc	ccaggccctc	cccgcctgat	gggtctctgt	ccgtccacgc	360
gggagacagc	gccacctgcc	ggtgagaagg	agcgttgctg	cgccggcacc	ageceagtee	420
tacgctcggg	gctcctgcag	gcctgggaag	gagggagggc	gcagctagaa	ggaagtctcg	480
cctgcccttg	cttccccgtc	tgtcagagtg	cctcgcatgc	aggeetgeet	ageggeettg	540

atcatgctct	ccctgtcacg	gaagtagaat	gtagtcaagt	ttttggactc	caagccattc	600
ttacaaaatt	gcgtcagagt	ggggattgta	ttataagaat	tgccactgaa	gagcagcgag	660
tggctgaaac	ctctgtgtgg	ctgccagtca	gcccctcccc	ggtgactgga	tcagcgaaga	720
atccagaagc	gaggttgcga	ggctgcagcc	cttggcatgg	ggagtccgtg	ggctgggcag	780
cactgcctca	gccgctggcc	tttcctgagc	agagtctagg	ctaagcggct	gttggaaata	840
gcagtagcac	ccggggcgag	accglgagcc	acagcggcgg	ccggagtctc	ccccagcccg	900
agctcaggcc	tgtgctggat	gcccaaagcc	tggcacagag	tttctttaac	cgcctttggg	960
aagtcgccgg	ccagtggcag	aagcaggtgc	cattggctgc	ccgggcctca	cagcggcagt	1020
ggctggtctc	catccacgcc	atccggaaca	ctcgccgcaa	gatggaggac	cggcacgtgt	1080
ccctcccttc	cttcaaccag	ctcttcggct	tgtctgaccc	tgtgaaccgc	gcctactttg	1140
ctgtgtttga	tggtcacgga	ggcgtggatg	ctgcgaggta	cgccgctgtc	cacgtgcaca	1200
ccaacgctgc	ccgccagcca	gagctgccca	cagaccetga	gggagccctc	agagaagcct	1260
tccggcgcac	cgaccagatg	tttctcagga	aagccaagcg	agagcggctg	cagagcggca	1320
ccacaggtgt	gtgtgcgctc	attgcaggag	cgaccctgca	cgtcgcctgg	ctcggggatt	1380
cccaggtcat	tttggtacag	cagggacagg	tggtgaagct	gatggagcca	cacagaccag	1440
aacggcagga	tgagaaggcg	cgcattgaag	cattgggtgg	ctttgtgtct	cacatggact	1500
gctggagagt	caacgggacc	ctggccgtct	ccagagccat	cggggatgtc	ttccagaagc	1560
cctacgtgtc	tggggaggcc	gatgcagctt	cccgggcgct	gacgggctcc	gaggactacc	1620
tgctgcttgc	ctgtgatggc	ttctttgacg	tegtacecca	ccaggaagtt	gttggcctgg	1680
tccagagcca	cctgaccagg	cagcagggca	gcgggctccg	tgtcgccgag	gagctggtgg	1740
ctgcggcccg	ggagcggggc	teccaegaca	acatcacggt	catggtggtc	ttcctcaggg	1800
acccccaaga	gctgcgggag	ggcgggaacc	agggagaagg	ggacccccag	gcagaaggga	1860
ggaggcagga	cttgccctcc	agcettecag	aacctgagac	ccaggeteca	ccaagaagct	1920
aggtggtttc	caggcccctg	ccctcccctt	ceteccatee	tigiccitci	ctccctcaga	1980
agcctcagga	cccaacaggt	ggcaggcagt	ggacagggtg	cccgccccac	agtgctttcc	2040
ccagcacccc	agagccagtc	gggacacccc	eegcageeca	tcctggtggc	tgtggaactg	2100
cactgggtgg	cgggcagatg	gtggaaggca	gcttaggaga	cctcaccaaa	gagaagatgg	2160
accggctctt	gctcccagct	cctattaggc	ccggggtggg	accagaggtc	ataggtgccc	2220
aacggcagcc	aaaccggcga	cgcacatgtg	tettttgttg	gtgtgtttgt	ttttttccag	2280
ggaggtctaa	ttccgaagca	gtattccagg	ttttctcttt	gttttatcag	tgccaagatg	2340
accigiigig	tcatataatt	taagcagagc	ttagcattta	ttttattctt	tagaaaactt	2400
aagtatttac	ttttttaaag	ctatttttca	aggaaccttt	tttigcagta	ttattgaatt	2460
tattttctaa	atcaggattg	aaacaggaac	itttccaggt	ggiğttaata	agccattcaa	2520
gtgccttaca	cagctttgaa	gaaactagga	ctgcagtggg	ctcggatagg	cccattgagg	2580
tttttagaaa	agcaggattt	gttttgttag	ggaggcatga	ttttggtgag	atctttctgg	2640

```
aagagttttc cgcctctttg tgatgctgaa caccccaag gttctcccct cccccgctg
                                                                    2700
cccaggtgac tggcaggagc tgcgactgcc acgtagtggt gcctgggccc gacagcgggg
                                                                    2760
                                                                    2820
ctetgggcat coogggtgac cttggcccat ctgcctgcat teccaeeee ttgggcctgg
ctggatccca ggcagaggga ccttgctgct gtgtgattgg aacattccca aatatcttgt
                                                                    2880
gaatttgtaa teaaattggt eteattggga aagaetetta attaagagge teaggeaage
                                                                    2940
                                                                    3000
acagaggcag cccgtgggtc tctgtctcag tctggaggca gcagggatgc tgctgggagt
                                                                    3060
ccatggcaca ggccacagcc cctcaccttg ccgcggtggc tggcagcacg cctgccttgc
tetgeeccat gecetgaaca ggeatgagag etceaegtee ectagtgeae eetgagaggg
                                                                    3120
ggctcacaag tgaccgatcc tgggtgcctc agggagctca ctgagggcgt gcaaagttga
                                                                    3180
aagtggcaag gctgggggag ggtgtcgggt agagggaaga gggcaggggg ctaggggagg
                                                                    3240
acteagagge catetgeagg gecaageeac aggaaggget gagetggagg tgggeaggge
                                                                    3300
tgctccaggc aggtcagagc agtgcagggg gaggaggga gaaagggagg aagctgggct
                                                                    3360
gtgtggtccc catgaaggca ttcagagtcc acctgcagac agcgagagcc ccaggaaggt
                                                                    3420
ttgcacaget gtgccccaag cacciiggee tecteteage tegcegagga ggcaegetag
                                                                    3480
agccgccttc ccggtgggag ccctctgtcc cacagggagc ggggagccag ctttgctggg
                                                                    3540
                                                                    3600
gecetacetg caigeceage ellacecele alteleacag cacagatgag gligagacea
tgcagtcaat gcattgctta aggtctctta titacaaaaa aaaaccitaa acatagtcgc
                                                                    3660
                                                                    3720
tgtcattcag acattcagag aaiggiigge cacaaacaai gaccaagiai igciiggcii
aacttgaagg cotgotgtot cottotgggg gtoagggaog cagotocaco otoacoacta
                                                                    3780
geceacectg ecegtgggea taacettgae gaagagagag aatgattgge atetgettit
                                                                    3840
ctcttttctt tgctaataat tctgttcctg gctgccgaga gtgaagtttc accatgtgga
                                                                    3900
ggtttggctc ctatcacctg gtggtctgat tcatacccta gcctgaggct ccactggaag
                                                                    3960
atttegeage cteagtgtat gggaaaceet tteeceagge ttgteecage actgeegete
                                                                    4020
                                                                    4080
eccaccetg agecaggate ccagaggatg gecatgeece gtgeetggea gaggtetggt
gecageactg ggagetgete egecetigee tiggggeega gggageeete giecaeeect
                                                                    4140
gcacagcagc tgggcacaga ggagcgctct tccatcttga ccaggactgc accaagaagc
                                                                    4200
accaggigte ticagecie aaccieeggg gegaecitei etteeageea eagieeeatg
                                                                    4260
agggeeecta gecagggaca etggtetgta aattgtaate ettteteeag eccagetete
                                                                    4320
cactigitee tigigigage igageaggea gigeaecici gagigieeet iitgiaagge
                                                                    4380
ccaggggttg cactgagtct gcagaggccg cgacctccta gaacgctgtg ggtgcaggtg
                                                                    4440
agccggcgtg tcctggggag atgctgccag cacacagggg ccctcctgct gccagcaggt
                                                                    4500
tggggtggtt aagtettatt agtgtetatt ettaaaatta agtgggetgg agaagaatgg
                                                                   4560
agetecacat gecageaceg latalggaat acaaaagelg gggaageagg geelgeelta
                                                                   4620
caggtgtggc tgactctgag cccaggcctg caggggtgga gggcagtccc tcagaatccc
                                                                   4680
agaggcagte ecageeteag aaceeaggat aggaaatggg tgtgtttagt ggggaaaggg
                                                                    4740
acggggtgca gacggcaggg ccagtatggg gcccctccc tetectotec tetectatgg
                                                                   4800
```

tgagcccagc	gtgggcaccg	ggccgtctca	gccatgttcc	cagggctggg	aggacagctc	4860
tggcccttct	taggcctagc	ctcgtcccaa	gctaaatgta	agccagttgg	gctgtgttaa	4920
aggaagcagt	gtttttggtt	cgattctgcc	tctgtagctc	aaggggggca	gcccccagag	4980
tcctgtgcat	tctgccaagg	ctccatagct	ttgccaaatg	cacggagete	tgccattccg	5040
gtgcagtgca	ggccttgcga	agggtttatc	tgcgttcgtc	tcggtgggct	tctcctgcat	5100
gggagttgtg	ttcctgtgca	agggggagct	ttgctccagg	acaggatgac	tgtcttccct	5160
attcttaggg	acaagtccca	agatgccaga	aaggcagtct	cccaaggacc	caccatgcag	5220
aagtgtcaat	aaaccacaag	ttctg				5245

<211> 4148

<212> DNA

<213> Homo sapiens

```
aaagatgtcc tcccctgatg ccacatcctg ttccaatgat cacgccttct ccagttccct
                                                                      60
tcacagccaa gcttctggag agcagcttct gcacaagctg tcttctattc ctctgttccc
                                                                     120
atccatgttc cagtccattc caggetggct cccatcctga tigcctcaca gaaactgttc
                                                                     180
                                                                     240
tttgcaggtc cccagccaag tecttattgc cacciccagc agectettic tgtcccacc
                                                                     300
ccettggace igicagcage allegaggea accgaeagea eligetgage igeleteeig
                                                                     360
teatggetgg acaegtggtg etgggeagge ttgeetggtg aggtgtggge aagetggaet
ccgtcttctt catccagtgc ctctggtctt aggcctgggt gtttgtctcc tctctgtgaa
                                                                     420
getetatgea gaactgigee aaggealegt ggacatagee atticeagig teticeeace
                                                                     480
cccagatgtg gagcaaccic agacccagcc agcigcciic aicaagcigi gacagagggc
                                                                     540
actecetget geettggaaa aageaegggg teetgeteea gggaatggtg aaatgaetgg
                                                                     600
attgctcttt atccagccca cagcagggga aagaaaggca actcgcaaag atgagatgga
                                                                     660
agaaggcacg tgagcagagg aggcagctcc caaagagagg gctgctcagg gggcttccca
                                                                     720
ggtgtagete teageagtge tgttgagaet tttgaaaaea actttggtae acaaaggeag
                                                                     780
ctttgtgage agageteett ecceteleee egggaaegge agggeaetgg gaeetetggt
                                                                     840
eggigeetee cacceactge agecetagig ceitagetee atgecegget geagececae
                                                                     900
tgctctggac tatggattgg acgtcagagc atattggagg ttgcctgtgt gttccccacc
                                                                     960
catecetteg glaacactet gecacactaa getetgtaca ageatgeace aacagteett
                                                                    1020
agittligige igigeaeigg ecieleggea aaggiggill eceleateae eticeigaig
                                                                    1080
gtgtttggtc agtcacctgt cagggtttgt gegggttggg ccccaaaaca gcatatgctg
                                                                    1140
ctctaagtct gctctctgca tgttttagaa acaaagtggc aagtctgccc tgaacctgta
                                                                    1200
```

agcatcaaat	aagcatgaga	gagaaaaaaa	catgatatat	tgctttactt	aataggttga	1260
atatggtagg	tctttgaaaa	tatgatgatt	caattttctc	aattttcttt	gctttaacca	1320
aaattctaaa	tgcagttttg	cctagttccc	ttttttttc	tttttttact	tttttttaaa	1380
cgtttgtaaa	aacctctttg	aggatgagga	gtcagtaaaa	ttccactccc	caagtggccc	1440
tgccccagac	aaaggttgct	ttccccttt	tigitettt	tatgccccga	agcactttct	1500
gcagtagcta	gagggacagg	tttccttcca	ggaaggattc	gagttcctgt	gcctgtgggt	1560
attaggagag	tatatatcct	gcctgaatgg	ggaagtcttc	taaaatggga	aagaagtggt	1620
ttcatctcca	cacagtgtct	tgtaaatctc	aacaaatgtg	tactgttaga	agtggcttcc	1680
gcttactgga	ttaactaata	ctttataggc	ttttcaggag	gccacatcac	tagcagtagg	1740
gagaacaaga	tgtcatttgt	gttcagtgta	agctgagtaa	acaggccctt	cctagagtgt	1800
cctggaaatc	acagcaaccc	attgaaaact	gccctcccca	ccagaacgtg	ctacgttctt	1860
tcttcatgcc	tatgtgtgct	ccattcctca	tttctacttg	gctcaagaaa	acatticige	1920
agtcaggtga	gacttttaca	aaagaggaga	aaatcaatgc	ctccttgaac	atgatgagat	1980
gtgagaactt	acaatgaaaa	aggcaataat	gatagaaatt	atttcttagg	tacagcaata	2040
gttgatagga	tgtgagggtg	ttaccttggg	gtgaagtgga	gaaggtccca	ggtgaattgg	2100
ctctcatgga	aatttggaat	tacgaaataa	acgtcctggg	ggttacccag	aatacagatt	2160
taaaagtttg	cctgtagagc	aaaataaaac	agtcagttgt	agtcattaat	ccttgaggcc	2220
caacgcagcc	gatgggttgg	tgtttgggaa	attctgagat	gggagtgaga	tctgatcgga	2280
tcctgggaag	atgtataccc	agttagaacg	tgtagggttc	tgggtccctg	gcaagtctag	2340
gtgggcgggt	gacagggaaa	gcatgggcat	ttttgtattg	ctgtcacatg	ctaacagagg	2400
tttgtaatta	tcttttggac	ccaaattata	gagacattca	cgagttttct	agccctcaca	2460
gtaacagagc	taagaattca	gatgtcagga	agtctgtgaa	tcttgatgga	ttttctgaga	2520
aacctgactc	aatggcatat	ataagaggga	agtaagactt	ttaagaaaag	aaaaagttat	2580
gcctcattcc	tcatgtggct	tccaataagt	atcttaggaa	cttatttcct	ttttaaaaaaa	2640
tatttttaa	attttaaaa	tttgatttta	aatttcaaat	aaatttaaat	aaattttaaa	2700
taaatttaa	ataaaatttt	acagagacgt	ggtctcacta	tgttgcccag	gctggattgc	2760
agtggctatt	cgcagttgta	atcatagcac	actgcagcct	cgaatttctg	ggcttgagca	2820
gtcctcccgt	ctcagcctcc	tgagtagctg	agactacagg	tgcacaccac	caageetgge	2880
tttatgtatt	tatttctgtt	catgcggaat	gattggttca	gaactgttcc	tttcccttcc	2940
atgatgtcct	tgacacagaa	ggttatgcct	ggctcccagt	caggcttcat	acttttggtc	3000
catgtaagtg	ctacccgttg	ctgggggagg	agtcatggtt	tatttggaaa	tgtcagttgc	3060
aatcatggtt	ctgtcatttg	actgcacagt	atcagaggag	cctgttaacc	tetetgtgee	3120
ttagtttctt	agcccatgaa	agagatcatt	gcctgaccca	gggactacct	caagggcttt	3180
tgatgaggac	aagtgacagt	aggaagatgc	aagagccttt	agtaccaagg	ttctcaacac	3240
tgactacatg	ctggaatgac	tgtgaagctt	ttaaaaaaatg	ttagtgccca	ctcttcccct	3300
gtaccccgg	acagttaaat	cagaacctca	gacagcaata	tgccttgaga	tgccttgaac	3360

catgcttgag	aaggaaggac	aaacacatta	ttatcttgga	agaattgcat	aaggcttatg	3420
acttaaaaaa	aaaaattctt	tttggaaaca	caagcatttc	tttaaggatg	accggatgtt	3480
gccgtatgta	tttatggcac	aagcaggtgt	tgtctaagca	gtttctctgt	ttgcttgtca	3540
tagcagcatt	tggaaactca	aacatgcttt	catttacata	aatagtttat	gaagctttga	3600
caacaaatgt	aaacagacac	gaaattataa	atctgctaaa	tatgtattaa	gggtattaat	3660
tattgaaagt	ccctttcccc	aaaactcaac	tcctatggca	attatgaact	ccattttacc	3720
aagaacattt	aagtgcctca	gcatctgtat	gatatagtgg	agcaggtgct	gacataggta	3780
ccagctgaca	tgatgtgtca	ctagctctgt	gggatgattg	ccacatacat	ggaacacctg	3840
ggagtgctgg	aaatgtactg	ggatcgaagt	gacaaagtgt	gttttcattc	acagtggagg	3900
ctacatcaag	caaggggagg	tccagccctc	ttgcaagtgt	ggtgagaggc	tctactagca	3960
aagacatggg	caccggagta	ggtcccgtgt	agcatgcggg	tgctgtagag	aaaattcagt	4020
gacgtacatg	gctctggttc	tggacacaaa	atctgtactg	gagaggaaat	gactgctgaa	4080
ataaggcgat	tgtatgaata	tttaaaatgc	ctggaacact	aaagtaaagt	aatgatattt	4140
caagtgtt						4148

<211> 2573

<212> DNA

<213> Homo sapiens

```
totgotgotc cgcgtgtggt aggagctacc agtctggggt ccgggctggg cgcattcatg
                                                                     60
atgcctgcct ggggtctgag caagtcctcc ccacggggtc tgagcaagtc ctccccacgg
                                                                    120
ggtctgagca aatcctcccc acggggtctg agcatgtcct ccccacgggg tctgagcaaa
                                                                    180
tectecceae ggggtetgag caagteetee ecatggggee tgagcaaate eteceeaegg
                                                                    240
ggtctgagca tgtcctcccc acggggtctg agcaaatcct ccccacgggg tctgagcaaa
                                                                    300
tectecceat ggggtetgag caaateette etatgeegte tgagcaagte eteceeatgg
                                                                    360
gitcigagea igicciccc acagggicig agcaagicci ccccacgggg icigageaag
                                                                    420
tecteccae ggggtetgag catgteetee ceaeggggte tgageaagte etecteeca
                                                                    480
cggggtctga tcatgtcctc cccacggggt ctgagcatgt cctctccacg gggtctgagc
                                                                    540
aagteeteec catggggtet gageatgtee teeceaeggg gtetgageaa gteeteecea
                                                                    600
eggggtetat gteeteeca eggggtetga geatgteete eecatgggtt etgageaagt
                                                                    660
cotcoccatg gggtotgage aagtootooc cacggggtot gagcaaatco tocccatggg
                                                                    720
gictgageaa alcettecta tgccgtctga gcacalcete cccaagetgt gaccgagtgt
                                                                    780
ccclcctgca ggtggaggat gttgctagga tgcaccttga aggcacccca gcctcgccgg
                                                                    840
```

```
agegeeecct cetegtagee tggggtgtgg etgggtggte tggggteetg ggtgeettgt
                                                                     900
gatgctggcc ccagggtcca ctcagcaccg tcctggtgtc gtcatcagct ggaggcttcc
                                                                     960
cggggcctgt gctgggggtg gagagcaggg agaggcagca gggttctcct cagggtgggg
                                                                    1020
tcgctgggaa gcaccatccc acctgtcaga ctggccttga ctgtagacac cccaggtgac
                                                                    1080
ctggaaggac agacggaccc caggtgatga gaaaggacca gagtctgacc tctcacccct
                                                                    1140
cetaagetet gaacteegt tggettgeet gaceteeaag teeteetggg getgaaceet
                                                                    1200
                                                                    1260
ctacagatgc ccctcctggg ccctggggtg ggcccggttt agctctccat tgtggctgaa
                                                                    1320
gcccccgggg cttcagtgct ggcttgaaga gggggtgggg ctccccaggc ctggggattg
gcagtttttt cctccctct tcccaaactt tcagactgga ccacttaaga ataatgaggt
                                                                    1380
ccaggtggtt ccgcttgagc ctggatcctc actggctgtg ggactgagct tcccctgccg
                                                                    1440
gtcccacctc ccaccgggag cagctaatga cagccagagg ctggaaggtg aagctccct
                                                                    1500
tggctgtcag gcgggccgca gggcagggc tgggcaggcc aagggcgcca ctctcctgcc
                                                                    1560
caggccaggg cacccgatca ctgcaccaca ccccttgtgg ccgtctgtcc agccagggcc
                                                                    1620
cigcigcagg igcticccgi gggacigtag ggagaacaat caagactici gccicciigg
                                                                    1680
tegageaggg etgeeteeee ateteateta etggeaagga ggetgggeae etteagggag
                                                                    1740
cttcagtttg ggaagaggga ggaggtctga ggtggatggt ggcgatggct gcgcagcagt
                                                                    1800
gagaatggac tgagtgccac tgatgtgtg gctccatggc tccgtggctc cgtggctccg
                                                                    1860
tggctcagtg gctcaatggc tataatggct agttttgtta catattttca ccataataaa
                                                                    1920
acaaaacatg tccaaggtgc tacaaggagg gaggagcccc tggagcaccc gcctgccatc
                                                                    1980
teceatetge caggeageat ecctecactg getetetggg aggggttega ggeeteeage
                                                                    2040
ctecetgtgg ecceeatetg cetecaggag attigtteec teteteetge eccgaaacee
                                                                   2100
tegaggeage cetgetettg gteactgeag aggaagtgge ceaggettgg eecaggeeag
                                                                    2160
ctgtggcctc cggaggcaag atgtggggac tcacagtgtt cgaaggccac accccccga
                                                                    2220
gcacatgggc tccagtgcct ctgaggcaaa gagcaggcag caccgtgcgc acagcagtgg
                                                                   2280
gagacacage acagecacea gggeageeee caggeagaeg gegggeetag agagggeggg
                                                                    2340
atgacacaag aaaggttete ettiggagae ggegaggtea ggeaggtggg agagggtica
                                                                   2400
cggtgcttga ggtgcagaga gaggatggtg gaatggaaaa cgtagggtga citgtcgggg
                                                                    2460
acaggeecag ggeeacaact egggeaggee tattgeeega gttttgggte ceateetgge
                                                                   2520
aggcagggga gagaattctg aatttttaa tgaaacggat agttgagggc tgg
                                                                    2573
```

<211> 2563

<212> DNA

<213> Homo sapiens

gtgaaatgtt	aggctttgtt	gatgaatgtc	atgaagagaa	tatgtacctt	tctgttgcct	60
tcacactcta	cctctggccc	tctgtgctgt	tcaaatgccc	atcttcctgc	tacctcctct	120
accttgaaac	attgcagggc	ttggagggag	cttgttctaa	agtctaagaa	gagctagatg	180
atttgtaaaa	ctttcttcag	accagctgcc	actgacagcc	tgcccggagc	cggacatggg	240
gcaggatcgt	gccgggattg	ctgtgactgg	atggtggaaa	attttgcaga	aacatctgtt	300
ctgtttggag	ggtccaaata	gttttaaaaa	catgtgctta	gccaaagctc	atatttcaca	360
aaacctttgc	aaatatctag	aagcttttct	tcttttctat	gtggacgtgg	aagcaaagga	420
gaggaaaatg	tggccacatg	tatgttttca	acttcttatt	tccaagtatt	tggctttttc	480
agggatgaga	accaatcaga	tcactttcgt	gaggtatgca	gtgcctctag	actgttctct	540
ctcttttgga	tagatacatg	aagtcttgaa	gaaagaaaaa	tttctgtaaa	cacaatggga	600
gagattacag	taatgctatc	aagctgtagt	tttaattgct	tgaaaataaa	cgaagaaaaa	660
ggttcacagc	tgtttgagag	tgaggaccaa	tcaagggcag	agcaacaaaa	aagctcccct	720
ttcctgggat	gactgccagg	actcagctct	ccacatctga	agacgtttta	caaagtgcag	780
tgtgccgtga	gcagggagag	aaaggcatcc	agagaaggcg	cgggaggact	tgagtgagga	840
gccaggtcct	ggcttcatcc	cagtctgtgg	gcctcaaggt	caggggagta	acgagctcat	900
ggccgacaga	ccgggatgac	agggacttct	taggggacaa	gtatgagttt	gttcaaactt	960
gggggcatga	gtttttgaga	acacggctca	acactcagca	tggtgaatgc	tgcagaccta	1020
gcatggagcc	gtacctggca	cctccaggag	aaaaaagcgc	cccaaactct	aaagctaaag	1080
gcctctgcac	atgattgcct	gtgaaccaga	gggttggaga	ttagttttct	cccccttag	1140
gtcattatgt	atgttccaag	ttgggcatgg	agagcagctc	ttctgccctt	tgaacctggt	1200
acagacccag	gaaacctggg	cctctccctc	ggtacctctc	attacaggtg	catggctcag	1260
gctcatggaa	caaatcagct	gacttttcct	ttgtttctta	atgctaggga	gcaggcaggg	1320
agctaaaggc	tgaaggaagt	tgaggcagtt	gtccttaaga	ctatctttag	tgaagtgaaa	1380
ggtgcagaat	ctgccatttg	tcatgtcacc	ttagaacaag	gcaaaatccc	cagggtacag	1440
acatccaatt	gatgtaccat	acttgatctc	caggttaaaa	tataatacag	ctatgatgca	1500
tgagtctcat	tgtgaaaaca	gctgattggt	gaggaaggtc	agtictcact	aaattggaga	1560
gatgaggccg	tgagatcaag	aggaagcagc	gctgagctgg	gagtccagat	agctggctct	1620
gctctctgct	ctgccaccag	ctgtggtgct	ggttaagtta	ctgggctctt	ccatcccctc	1680
tctgccttgt	cagtaggcag	attggatgat	gtgtaagttc	ctcctgtgct	gaagatcctt	1740
gaactgagga	cctgatttcc	agagcccagg	gaacatctta	gaaatggagt	aaattacatg	1800
agattttccc	aggggaggcc	ttgatcacat	tttgtacaac	attcagtcat	gtatggttgc	1860
tatgatacca	ggcagcattt	tgaaaccata	cacagggatg	agtctttcag	tcagtggcct	1920
aaaccatctc	cctttgctgc	agagccagct	tttctgcaat	tccaggggaa	agtatgggca	1980
attgttaata	ccccaaagat	tttatatgat	tttaaaacaa	agtggccaac	agtgtcaaca	2040
ttgtttacca	gtgactcgtg	tcttttttt	cctttgtcct	cctccttttt	taaaaaataa	2100

```
2160
acctataatg taaacctctc aatttgcttt acttttcctg ctcttgagat tttcatgtgg
                                                           2220
ccctgattaa aattttaatt tgtcagtaga gtcaaatctt attagtgcca ttccagcaat
                                                           2280
                                                           2340
tgggcactgg gatcatttgc aaggtcttca gggaagtttg cctttgcaca gtttaggaaa
gatictgtta attaggtgaa tggtataatt gatacgacaa gaggattgtt taacttaagg
                                                           2400
gaagcaattt attatgcatg catgagaagc ttctaggtat ttactgacca attgcatgcc
                                                           2460
cattacatat cctttttgta ttttagagat aataatcatc ttatattgtt tacctcctag
                                                           2520
cccagttttt ggcacacttg aaagtactac aaattgtctt tat
                                                           2563
```

<211> 3219

<212> DNA

<213> Homo sapiens

<400> 2068

catcagtaaa ggcacggagg tgggaaacta tgtagtgtgc aaaggaaaag tcagatgatg 60 120 gtgatgataa tggagagact gacagcagca gacatctttt tgagcactta gtgtgttcca ggtgtgtgta ccaagcacta tcctggctga atctcatcag attggatggc aggaagtaaa 180 acticagagi ccatgittee aatgeegeag ctaccetgie teteatgaat gaggagetgg 240 aggagettgg attegttgea gtttttttt tttttttett tttttttea ggaeggagte 300 tegetetgte geecaggetg gagtgeaatg gtgegatete ggeteaetga aaceteeace 360 420 tectgggtte atgegattet catgeeteag eeteecaagt agetgggate acaggtgeee accaccacge eeggetaatt titgtatitt tagtagagae ggggtticae cataitggee 480 aggetggtet egaacteetg aceteaggtg atecacecae eteggeetae caaagtgetg 540 600 ggatlacagg cgtgagccac catgcccagc cagattettt gcagtttaac acgtttccag agagtgtgtt ctaggtcagg ccctggtgct ggaagcaggg acccatgagg gccaaggcct 660 ggtccttgcc ctcaaagget gacccagtta tagtccaggg tggtgagggg ccagctgggg 720 780 ctgctcatag cctctggcag ccaaagtggg gtattgaggg gctggggagg aagcgttgtg gigggggggc cigcagicci aggcagggia giatgaggcc cagcitcati gcicagiagi 840 cacatcatet caggeaagee actiggeele tetgageete agtigeetet geteagaagt 900 aacaacctga actiggacta teagggaage eeagggeeca eagetiggte etaggaaggg 960 ctlagcaaac gggggtggtt gtccttcttg gaagccacat ttgtttgcct ggtgagtggt 1020 ggagggcact gctaggcctg ctagggctga cacggccaga gtcagatgac ctcatctcac 1080 alccagcagg tgaaatgcag tetttgatee ettgaaacce accetetagg accaaggtea 1140 ctgcagtatt ggataggacc tcagggagtt agcagggggc tcatggttaa gagtgtgaac 1200

tacggcttag	acctacaggg	ttccctgccc	agctcctcca	caaaccagct	gtgcaaccct	1260
agacaagtga	gttaatgtcc	ctgggcctca	gtttcttctt	agtaaaatgt	gtgtagccat	1320
agagggctgt	tatgaggatt	cagtcaaatg	acacatgatg	tcttgggcac	acctggcgtg	1380
gattatggcg	cctgtaggag	caggagggct	tcctggagga	gggggctagt	tgaacagagt	1440
ctagaaagta	tagattggga	agagcactct	gggaggcagg	atcaccatgt	gcaaaggctc	1500
agagaatgcc	acccactacc	tcctggaaat	caaggggatt	ctgtgtgtcc	aagggcattg	1560
gtggtctcta	ggcccccgac	ctgtgtctgg	gaggtgtcaa	ggggaagcca	gatccgaggc	1620
ccacacttgc	atgttttcag	gtgaggtcca	gagatatatc	cagagaggag	tggaagggct	1680
cggagaccta	cagccccaat	actgcatatg	gtaaggcccc	agctctgagc	ccacctgcag	1740
gagcttcagc	ccttgggccc	agcctccaca	tgaccctccc	atatcccagc	catggcattc	1800
tggctgggaa	gccttctctt	ctgccctgc	ctagagggtt	ggggagcaca	tgggccccta	1860
gagagggagg	gacacctcgc	tggtacaggg	atgtgagtgc	agaccctgcc	atcccatcct	1920
acaggtgtgg	acttcctggt	gcccgtgatg	ggctatatct	gccgcatctg	ccacaagttc	1980
tatcacagca	actcaggggc	acagcictcc	cactgcaagt	ccctgggcca	ctttgagaac	2040
ctgcaggtga	gccggacatc	ctgccctgtc	ctccctggc	cacagactta	gtcttaatcc	2100
aagctgattc	gggtggctag	tggccactcc	ctcttgtgca	gggcctcaat	ccccaggcac	2160
caccctgca	ccaacaggga	gagaattaga	gctggggtgg	ggttgggccc	ttattgttca	2220
aggggatgct	gagtgccagg	ctgttagctc	cagagacggc	ccagagaggc	cgagtgcatc	2280
acgcagggtc	acagagcaca	ctaatactgt	ctcagccaga	gctggggaag	tagctgctgg	2340
ccaggagcat	accatgtagg	gaggagaccc	tgaccttacc	tgcaccttct	gtatccagaa	2400
atacaaggcg	gccaagaacc	ccagccccac	cacccgacct	gtgagccgcc	ggtgcgcaat	2460
caacgcccgg	aacgctttga	cagccctgtt	cacctccagc	ggccgcccac	cctcccagcc	2520
caacacccag	gacaaaacac	ccagcaaggt	gacggctcga	ccctcccggc	ccccactacc	2580
teggegetea	acccgcctca	aaacctgata	gagggacctc	cctgtccctg	gcctgcctgg	2640
gtccagatct	gctaatgctt	tttaggagtc	tgcctggaaa	ctttgacatg	gttcatgttt	2700
ttactcaaaa	tccaataaaa	caaggtagtt	tggctgtgca	gttcccacca	gtacttctgt	2760
ctgggtggat	aggggaaggg	gggcacccca	gccaactctc	agccagcacc	cagcctctct	2820
gggccatgtg	gtggcagaaa	cagaaggcca	gacaggctcc	ctgggaacca	gggactctgg	2880
atcatgaggc	acttcacctg	tctgaacttg	ggtttccctc	ttttaaaaaa	atttttaggc	2940
ggggcgtggt	ggctcacacc	tgtaatccca	gcactttggg	aggctgagac	gggtggatca	3000
cctgaggtca	ggagttcaaa	accggcctgg	ccaacatggc	aaaaccgtct	ccactaagaa	3060
atacaaaaat	tggctgggtg	ttgtggcggg	cgcctgtgat	cccagctact	cgggaggctg	3120
aggcagggag	ggttgcttga	gcccgggagg	tggaggttgc	agtgagccga	gatcgtgcct	3180
gtgcactcca	gcttgggcga	aggagtgaga	ctccatctc			3219

```
<210> 2069
<211> 3341
<212> DNA
<213> Homo sapiens
```

						1.007 2000
60	ttgccagggc	ggcgcggcgg	gcgggtgagg	tcagaaagga	caccacagcg	gaacgaaaac
120	ggaagtgggg	acagtggcga	tcaactagtg	ggctgatgag	cgtcgggcag	atcttcttag
180	atctctgcag	aagggcgctg	gagctaggaa	aggctgaagg	agcgagagga	gcgctgagca
240	gagggcaggg	atggggtccc	gccagaagag	ccggaggaag	cttttgtctc	cctgggaggg
300	gaacggggag	agcccacaga	tgtcattttg	ggagcatgcc	aagaaaacga	ctcacacagc
360	ttaaacagtt	gccgaggcgg	ttattcggta	ctgctcgggc	ggaggaccgg	cggagccact
420	gggcagcacc	cagggttgct	gtgcagtctc	ctggagcagg	ccagccggga	cagggctgga
480	cacattgttc	tccacacttg	acgggagctt	cgaataaact	gagcaccgaa	gagacccttt
540	tcaccctcag	tccttggcag	tagcgctcat	ttcctgatgc	cagacgcagg	ccgcgagttg
600	gcaagtaggt	gattccaggt	tgaatgcttg	accttcagga	agttgccgtg	tgaactacac
660	caggaaagat	ctcagaaact	ctgaaggtcc	cctccagtca	agcttcctcc	actggagggg
720	ctgtgaccct	cagtctgttt	cctgaccacc	tatttctact	cctagaaaat	gatgaaagag
780	caggctctgc	ctttaatacc	taagatctgg	cagtaagtca	gaacagtgtt	ttgtagctgc
840	tgttttctcc	tctctgaacc	tcacttaaac	catgggcaag	tggtgtgagt	cacttgctag
900	taaatgacga	atgcacgttg	gtigtagiga	acctcccagg	ctgaggtaat	ttttttaaaa
960	atgictggaa	accttgcata	gattccccac	ccactagctg	tcatccttta	gctacattcc
1020	acttacttcc	acagttgtgc	tgaatgaagg	ttctcttgta	ctcagaaata	cattctggtg
1080	gctgagggcc	tttcaggttt	tcgtttgttc	gaggaatgtc	ttaactgaca	taaagtttca
1140	ccagagaagc	ttttgtcaag	taataaataa	tatcatagtc	ccttccaccg	ccagaaggct
1200	gatctgaaca	ggaaatgacg	agatagtggc	cttaaagaaa	agagacaagg	taacaaaggt
1260	agggagctac	ggaggaatcc	aacagctact	acccagcgga	acaagctcaa	agcatataaa
1320	ctgactaaca	tctggaatac	acagattctt	caggctgaaa	gttacttgtc	accgagaaaa
1380	caaaaaagtg	cagctattta	aggtatggaa	caacctgaga	gtacacagag	aaactgaaga
1440	atttcagtgc	tgcagaacaa	cctccagata	caagaatcag	acgaagaaga	gagagattga
1500	aagttgcagg	tttgaagcgg	tecaatecag	aaggaaaata	gctcttgcaa	ttaaaacagc
1560	acattacagg	agaaatacag	agcaggagaa	ttaaaggaaa	cattgctata	caatgaggga
1620	caggcccagc	acgggaagta	cttcaaagac	gctgagacag	gaaagtccaa	aggagacaaa
1680	aggctactgg	gccagacagg	aactgagcga	ctggagaaac	gaaaagatta	tectecagga
1740	gcagcaaagc	cttgaagttg	aggcccaggc	cttaatatga	aagaagagag	gaaagagaaa

```
ggtttatttt tgaatactcc tgtggcatca acagagagaa ccagcagttc aagaaggaat
                                                                   1800
                                                                    1860
tactgcaget aattgagcaa gcccagaaac taacggctac tcaaagccac ttagaaaaca
                                                                    1920
ggaagcagca gctgcagcag gaacagtggt atctggagtc cttaatccag gcgaggcaga
                                                                   1980
gactgcaagg aagtcataat cagtgcctaa atagacagga tgttccaaag accacaccca
gtetteecca aggeaccaaa teaaggatta ateeaaagta aettetaaaa taacaetgat
                                                                   2040
                                                                   2100
taaataagaa ctggagcaag tactcttaag tgctacatta acctggttag aaaggctgtt
ggattccaga ttgctattgt aaaatctcca tcatgatgtg ttggagtgaa ggattagatg
                                                                   2160
gttttatcca acagtcctac tagatatttg gtaaccagct tcccttaact agctttttct
                                                                   2220
                                                                   2280
ttaaatactc gttaataagc tattccacaa acctccagtt aacctaacac atgaccctaa
                                                                   2340
cctagccatt taccatacat caaactagct aaaggaaacc aacctaagga agtgaaaaca
gttgtgattt atttcatcta gctaaattgt atttctttat agagaaagta cctttaagga
                                                                   2400
tagcattcca aatagacttt gaatagcgtt ctgccagttt atcctcattc cttttgacca
                                                                   2460
acttagcaga caaaagcagt ttttacaagc tctttgtgag tttgtgccag tgaccaggta
                                                                   2520
                                                                   2580
gctccttcta gttttctcat gagtgaaaaa gcattctgat aacagcaagt ccagtaagtg
ctaggcagag tgacctttca tctgatgcta agcccctaca agtttgagaa ggtaagaaaa
                                                                    2640
                                                                   2700
gatgaaggag acatatatta ggtcagctct tacttttgaa aatgttttat ttgaagaaac
                                                                   2760
acctgtagca ttgaggtgac tgaatgcctc cacttatttc aggaaaacgt atccaaaaaa
                                                                   2820
agttgaaata tttggacaac ttttttttta agtgccatcg atttccctag cagcattcta
                                                                   2880
aaagatagca agtaaaatga tgtttgttat cctaaatgct ttagttttag gtcatttatt
                                                                    2940
aattttetta eaggtgeaet ttetagtaea tgaagtatee tttgtaatta atgtgtgeea
                                                                   3000
tatgittati eccattiagi ataaciataa attatattii aaaltatata tiittaggat
agttatattt ttttttgggt tctacgacat tgaagttgga ctagtgattt atttgaatgc
                                                                    3060
                                                                   3120
tgaatcctag tataggggaa tataatctta tattttaaca ggggtcctct atgggaaaat
aggatgaact ttgtttccca gaaattgtta agtgatgaaa aacttcaaaa taattttcct
                                                                   3180
gcattitcig cittattiac aigtaaagig aattccciga aaaitggati taaaaagcat
                                                                    3240
                                                                   3300
teteetteaa tgtgeettta eettggaget ttaacaactt ttetgttaaa tatgtagttt
tttattaaac aatgttatta aataaaaaca tttatccact g
                                                                    3341
```

<211> 2517

<212> DNA

<213> Homo sapiens

<400> 2070

aaaagaccca tgagacctct cctcgtctgt gcacagactg gtggccgact ctggagccca